Human macrophage migration inhibitory factor potentiates mesenchymal stromal cell efficacy in a clinically relevant model of allergic asthma

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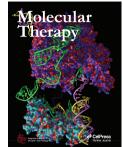
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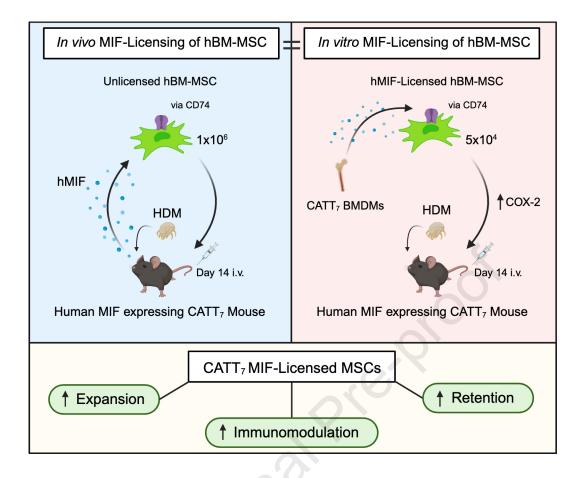
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English and colleagues discuss how mesenchymal stromal cells significantly attenuated house dust mite-induced airway inflammation and airway remodelling in high MIF expressing CATT₇ mice. Blockade of CD74 or COX-2 function in MSCs prior to administration attenuated the efficacy of MIF-licensed MSCs *in vivo*.

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21 MIF enhances MSC efficacy in asthma

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31 Abstract

Current asthma therapies focus on reducing symptoms but fail to restore existing structural 32 33 damage. Mesenchymal stromal cell (MSC) administration can ameliorate airway inflammation 34 and reverse airway remodelling. However, differences in patient disease microenvironments seem to influence MSC therapeutic effects. Polymorphic CATT tetranucleotide repeat at 35 36 position 794 of the human macrophage migration inhibitory factor (hMIF) gene has been 37 associated with increased susceptibility and severity of asthma. We investigated the efficacy of human MSCs in high vs low hMIF environments and the impact of MIF pre-licensing of 38 39 MSCs using humanised MIF mice in a clinically relevant house dust mite (HDM) model of 40 allergic asthma. MSCs significantly attenuated airway inflammation and airway remodelling 41 in high MIF expressing CATT₇ mice, but not in CATT₅ or wildtype littermates. Differences in 42 efficacy correlated with increased MSC retention in the lungs of CATT₇ mice. MIF licensing 43 potentiated MSC anti-inflammatory effects at a previously ineffective dose. Mechanistically, 44 MIF binding to CD74 expressed on MSCs leads to upregulation of COX-2 expression. 45 Blockade of CD74 or COX-2 function in MSCs prior to administration attenuated the efficacy 46 of MIF-licensed MSCs in vivo. These findings suggest that MSC administration may be more 47 efficacious in severe asthma patients with high MIF genotypes (CATT_{6/7/8}).

48

49 Introduction

Allergic asthma is characterised by chronic airway inflammation and airway remodelling 50 51 which refers to the structural changes in the airways. Currently, there is a heavy reliance on 52 inhaled corticosteroids and long acting β 2-adenoceptor agonists in the treatment of allergic asthma. The recent introduction of novel biologics such as benralizumab and dupilumab 53 54 targeting Th2 cytokine receptors and tezepelumab targeting the alarmin thymic stromal lymphopoietin (TSLP) have been shown to significantly reduce allergic airway inflammation 55 leading to reduced exacerbations and improved FEV₁ values.^{1,4} However, not all patients are 56 responders, and evidence for biologics to reverse existing airway remodelling in patients is thus 57 58 far limited.⁵ Thus, there is scope for novel therapeutics with capacity to attenuate inflammation 59 and reverse remodelling to address the pitfalls in the current treatment and management of 60 allergic asthma.

61 Mesenchymal stromal cells (MSCs) have immunomodulatory and anti-fibrotic properties and have proven therapeutic effects in a range of allergic airway inflammation models and are 62 currently under investigation in two clinical trials for asthma (NCT05147688, NCT05035862). 63 Administration of MSCs intratracheally or intravenously has been shown to be effective in 64 reducing airway inflammation and airway hyperresponsiveness in ovalbumin (OVA),⁶⁻¹⁷ 65 HDM,¹⁸⁻²⁴ and aspergillus hyphal extract^{25,26} models. However, other studies fail to 66 demonstrate efficacy in experimental asthma models.^{7,14,23,24,27,28} To understand the 67 mechanisms involved and to make MSCs a viable therapeutic in the clinic more focussed 68 translational work is needed. 69

70 Under basal conditions, for example in healthy animals or individuals, MSC administration 71 does not seem to alter immunological status or function (homeostasis is preserved). MSCs only become licensed to an anti-inflammatory phenotype in the presence of extrinsic factors.²⁹ Once 72 licensed, MSCs modulate their surrounding microenvironment.³⁰ Importantly, their therapeutic 73 effect is blunted in the presence of IFNγ, NF-κB or TNFα receptor blockade/inhibition.³¹⁻³³ 74 Moreover, in the absence of appropriate signals to license anti-inflammatory functions, MSCs 75 may even exacerbate disease.³⁴⁻³⁶ Licensing has been shown to improve MSC therapeutic 76 77 efficacy by activating MSC anti-inflammatory characteristics prior to administration. Licensing through exposure to hypoxia,^{37,38} inflammatory cytokines,^{39,40} and pharmacological 78 factors⁴¹ have all been shown to improve MSC efficacy in a range of inflammatory diseases. 79 Moreover, licensing of MSCs with serum from HDM-challenged mice¹⁸ or with serum from 80

ARDS patients⁴² enhanced MSC therapeutic efficacy *in vivo* in pre-clinical lung disease models. However, there are also *in vitro* studies reporting differential and in some cases negative effects of patient samples (acute respiratory distress syndrome (ARDS) versus cystic fibrosis (CF)) on MSC survival and function.⁴²⁻⁴⁴

85 Macrophage migration inhibitory factor (MIF) is an important regulator of host inflammatory responses demonstrated through its ability to promote the production of other inflammatory 86 87 mediators. For example, MIF has been shown to amplify the expression of TNF, IFNy, IL-1β, IL-2, IL-6 and IL-8 from immune cells.⁴⁵⁻⁴⁸ This augmentation of immune signals contributes 88 89 to MIF-mediated pathogenesis by acting to sustain inflammatory responses. This has been 90 shown in a range of inflammatory diseases where the absence of MIF is associated with lower 91 levels of pro-inflammatory cytokines resulting in reduced pathology. For example, MIF knockout (MIF^{-/-}) mice display a less severe phenotype when exposed to OVA compared to 92 control mice⁴⁹⁻⁵² and the use of anti-MIF antibodies or small molecule inhibitor (ISO-1) results 93 in reduced Th2 cytokines in models of allergic airway inflammation.^{51,53-56} High levels of MIF 94 as a result of longer CATT repeats such as CATT₇ have been shown to increase severity in a 95 range of diseases including severe anaemia (57), pneumococcal meningitis,⁵⁸ multiple 96 sclerosis,⁵⁹ tuberculosis,⁶⁰ and COVID-19.⁶¹ Importantly, associations between the CATT 97 polymorphism and asthma incidence and severity have been observed.⁵² Not only do these 98 99 studies show the pivotal role that MIF plays in pro-inflammatory diseases they also affirm the 100 importance of differences in the MIF CATT polymorphism.

101 Our previous work established a dominant role for MIF allelic variants in the severity of HDMinduced allergic asthma.⁶² Using humanised high-expressing and low-expressing MIF mice in 102 a HDM model of allergic airway inflammation we demonstrated the pivotal role MIF plays in 103 104 exacerbating asthma pathogenesis. High levels of human MIF resulted in a significant increase 105 in airway inflammation as a result of elevated levels of Th2 cytokines promoting infiltration of 106 eosinophils into the airways. Furthermore, high levels of MIF were associated with airway 107 remodelling with significant mucus hyperplasia, subepithelial collagen deposition, and airway hyperresponsiveness generating a more severe asthma phenotype. MIF has been shown to 108 promote MSC migration in vitro,⁶³ however, the effect of MIF on MSC immunosuppressive 109 110 function or therapeutic efficacy in vivo is unknown. Here, we sought to investigate the 111 relationship between MIF and MSCs in vivo and to define conditions for optimal MSC 112 therapeutic efficacy. The high MIF expressing CATT₇, low MIF expressing CATT₅, and WT

113 mice were used as a platform to investigate the role of MIF on MSC efficacy in a clinically

114 relevant HDM-induced mouse model of allergic airway inflammation.

115

116 **Results**

Human BM-MSCs significantly reduce airway remodelling in CATT₇ mice challenged with HDM

119 Firstly, to investigate the impact of high- and low-expressing MIF alleles on MSC treatment of 120 allergic airway inflammation we examined the lung histology. CATT₇, CATT₅, and WT were 121 randomized to HDM or mock (saline) intranasally 3 times a week for 3 weeks. Mice were then 122 further randomized to 1x10⁶ human BM-MSCs or equal volume saline administered via tail 123 vein injection on day 14. On day 21, lung tissue was removed, formalin fixed and sectioned 124 onto slides (Figure 1A). Slides were stained with PAS to highlight mucin production to assess 125 the level of goblet cell hyperplasia. CATT₇ mice exhibit significantly higher levels of goblet 126 cell hyperplasia compared to WT and CATT₅ mice. Administration of BM-MSCs reduced the 127 level of goblet cell hyperplasia in all groups to almost background levels, with a significant 128 reduction in the number of mucin secreting cells in the airways of HDM challenged CATT7 129 mice (Figure 1B, 1C).

130 Subepithelial fibrosis was significantly increased in HDM challenged CATT₇ mice compared to the lower MIF expressing CATT₅ and WT groups. BM-MSC administration reduced the 131 level of subepithelial fibrosis to almost background levels in all groups with significantly 132 133 reduced subepithelial collagen deposition in HDM challenged CATT₇ mice (Figure 1D, 1E). 134 In CATT₅ and WT mice challenged with HDM, BM-MSC administration had a small but not 135 significant therapeutic effect. BM-MSC administration significantly mitigated increased 136 inflammatory infiltrate and H&E pathological score in CATT₇ mice challenged with HDM 137 (Figure S1).

Human BM-MSCs significantly reduce airway inflammation in CATT₇ mice challenged with HDM

140 Total cell counts were significantly elevated in the BAL fluid of CATT₇ mice following HDM

141 challenge (Figure 2A). MSCs significantly reduced the number of total infiltrating cells in the

142 BALF of CATT₇ mice (Figure 2A). Differential cell counts identified eosinophils as the main cell infiltrating into the lung tissue following HDM challenge and MSCs significantly 143 144 decreased infiltrating eosinophils in CATT₇ mice yet had no effect in the CATT₅ and WT groups (Figure 2B). IL-4 and IL-13 were significantly elevated in the BAL fluid of CATT₇ 145 146 mice following HDM challenge (Figure 2C, 2D). These Th2 cytokines are not significantly 147 upregulated in CATT₅ or WT mice. While MSCs significantly decreased IL-4 and IL-13 in 148 CATT₇ mice, MSC treatment did not reduce and in some cases increased Th2 cytokines in the BAL fluid of CATT₅ and WT mice (Figure 2C, 2D). These data show that BM-MSCs are 149 150 effective at alleviating eosinophil infiltration and reducing Th2 cytokines in a high MIF expressing model of allergic asthma and that MSCs require a threshold level of inflammation 151 152 to mediate their therapeutic effects.

High levels of hMIF significantly enhance BM-MSC retention in a HDM model of allergic asthma

Next, we analysed the biodistribution of MSCs following administration into HDM challenged 155 WT, CATT₅ and CATT₇ mice. 1x10⁶ fluorescently labelled BM-MSCs were administered i.v. 156 via tail vein injection on day 14. On day 15 mice were sacrificed and the lungs prepared for 157 158 CryoViz imaging (Figure 3A-C). Significantly higher numbers of labelled MSCs were detected 159 in the lung of high MIF expressing CATT₇ mice compared to the low expressing CATT₅ or 160 WT littermate control (Figure 3C, 3D). However, the number of clusters of labelled BM-MSCs 161 within the lungs remained unchanged among the groups (Figure 3E). Taken together, these 162 data suggest prolonged MSC-pulmonary retention time increases the number of MSCs retained 163 at the site of inflammation 24 hr post administration. These data suggest that high levels of 164 MIF may provide a longer window for MSCs to carry out their therapeutic effects.

The influence of MIF on MSC expression of immunomodulatory factors and MSC cytokine licensing *in vitro*

167 MSCs mediate their therapeutic effects via expression or production of secreted factors *in vitro* 168 and *in vivo*⁶⁴ and licensing with proinflammatory cytokines such as IFN γ or TNF α ⁶⁵⁻⁶⁹ can 169 enhance expression of immunomodulatory mediators. Here, we characterised the effect of 170 recombinant human MIF on the expression of IDO, COX-2, PTGES, ICAM-1 and HGF in 171 untreated MSCs or MSCs licensed with IFN γ or TNF α . rhMIF (1ng/ml) stimulation alone did 172 not increase expression of IDO, COX-2, PTGES, ICAM-1 or HGF (Figure 4A-E) in human

6

173 BM-MSCs. Following licensing with $TNF\alpha$, MIF stimulation enhanced MSC expression of 174 COX-2 and PTGES (Figure 4B-C). In IFNy licensed MSCs, rhMIF stimulation did not enhance MSC expression of IDO or HGF and significantly reduced ICAM-1 expression (Figure 4A, 175 4D, 4E). We confirmed these findings at the protein level for IDO and COX-2 using 176 177 intracellular flow cytometry (Figure 4F-G). Using increasing doses of rhMIF (1, 10 or 100 178 ng/ml), we showed that COX-2 expression is increased in a dose-dependent manner by rhMIF 179 stimulation in TNFa licensed MSCs, with COX-2 expression plateauing at 10ng/ml of rhMIF 180 (Figure 4H). The MIF receptor CD74 is expressed by MSCs, however, rhMIF stimulation (dose 181 range 1, 10 or 100ng/ml) does not enhance CD74 expression (Figure 4I).

182 CATT₇ MIF licensing enhances MSC expansion and immunosuppressive function *in vitro*

In order to investigate the effect of endogenous MIF from CATT₇ mice on MSC expression of immunomodulatory factors, we generated bone marrow-derived macrophages (BMDMs) from CATT₇ mice and used the conditioned medium (CM) as a source of endogenous human MIF (Figure 5A) to license MSCs. The concentration of human MIF in CATT₇ BMDM CM ranged from ~3000–4000 pg/ml (Figure S2).

MIF may have a negative role in the regulation of IDO expression as MIF^{-/-} mice produce more 188 IDO,⁷⁰ however, MIF has an established role as an upstream positive regulator of 189 cyclooxygenase 2 (COX-2) through the activation of the MAPK signalling pathway.^{71,72} IDO. 190 191 COX-2 and PGE2 are widely reported mediators of MSC immunosuppression.^{39,73} MSCs constitutively express COX-2 but not IDO. IFNy licensing of MSCs leads to expression of IDO 192 while TNF α enhances MSC COX-2 expression.⁶⁸ Here we show that CATT₇ MIF stimulation 193 194 reduces MSC IDO production (Figure 5B-C), however, the percentage of COX-2 expressing 195 MSCs was significantly increased following CATT₇ MIF stimulation (Figure 5D-E). Human 196 MSCs express the MIF receptor CD74 and this expression is maintained and not increased following exposure to CATT₇ MIF CM (Figure 5F). In line with other studies,⁷⁴ we show that 197 198 IFNy stimulation leads to significantly increased MSC CD74 expression and CATT₇ MIF CM 199 does not significantly alter that (Figure 5F). This aligns with our data showing that MIF does 200 not enhance IFNy regulated IDO expression. Given the potentiating effect of MIF on the TNFa 201 regulated gene COX-2 in MSCs, we examined the influence of CATT₇ MIF on MSC 202 expression of the TNFα regulated genes TSG-6 and PTGS2. The presence of CATT₇ MIF CM significantly reduced TSG-6 in TNFa stimulated MSCs (Figure 5G), but did not significantly 203

alter the expression of PTGS2 (Figure 5H). MSCs licensed with high levels of hMIF from
CATT₇ CM displayed enhanced suppression of T cell proliferation compared to the untreated
MSCs, however this was not statistically significant in the frequency of proliferating CD3⁺ T
cells (Figure 5I) or the number of proliferating CD3⁺ T cells (Figure 5J). The presence of SCDabrogated the enhanced suppression mediated by hMIF licensed MSCs, as the number of
proliferating CD3⁺ T cells was significantly increased compared to the ^{CATT7}MSC group
(Figure 5J).

- Previous studies have shown that MIF has the ability to support cell proliferation *in vitro*.⁷⁵⁻⁷⁷
 Increasing the number of MSCs within the inflammatory niche could prove to be important in
 enhancing MSC immunoregulatory effects. High levels of MIF significantly enhanced MSC
 expansion *in vitro* compared to the complete medium control group (Figure 5K). Blockade of
 MIF using SCD-19 confirmed the role of MIF in driving MSC expansion (Figure 5K). This
 data might help to explain the enhanced retention of MSCs in CATT₇ HDM challenged mice
- 217 (Figure 3) but further experiments would be required to confirm that.

218 Titration of BM-MSC doses in CATT₇ mice challenged with HDM

Next, we investigated if MIF licensing could improve MSC efficacy in the high MIF expressing 219 220 CATT₇ mice challenged with HDM. To do this we first investigated the dose at which MSCs lose efficacy. MSCs at doses of 1×10^6 , 5×10^5 , 1×10^5 , and 5×10^4 were administered i.v. into 221 222 HDM challenged CATT₇ on day 14 (Figure 6A). MSCs maintained efficacy as low as 1x10⁵ 223 cells with reduced immune cell infiltration (Fig. 6B, 6C) and reduced Th2 cytokines IL-4 224 (Figure 6D) and IL-13 (Figure 6E). We observed that BM-MSCs were no longer able to carry out their immunosuppressive effects at a dose of 5×10^4 . At 5×10^4 BM-MSCs were unable to 225 226 reduce the number of eosinophils infiltrating into the lungs (Figure 6B, 6C) or regulate Th2 227 cytokine production (Figure 6D, 6E).

228 MIF licensing restores MSC efficacy at low doses in CATT₇ mice

To investigate the effect of MIF licensing on MSC therapeutic efficacy MSCs were first licensed *in vitro* by stimulation with bone marrow-derived macrophage (BMDM) conditioned media from CATT₇ or KO mice for 24 hr. 5x10⁴ MSCs, MIF licensed MSCs (^{CATT7}MSC), or MIF KO licensed MSCs (^{KO}MSC) were administered i.v. into CATT₇ mice via tail vein injection on day 14 in HDM challenged mice. On day 18, BAL fluid was collected, cell counts were performed and Th2 cytokines were measured (Figure 7A). Only ^{CATT7}MSC administration

significantly reduced total cell counts and number of eosinophils in CATT₇ mice challenged
with HDM (Figure 7B, 7C). ^{CATT7}MSCs markedly reduced IL-4 and IL-13 levels compared to
control group although not significantly (Figure 7D, 7E). The control MSC group and the
^{KO}MSC group displayed similar levels of immune cell infiltration and Th2 cytokine production
suggesting the effects observed in the ^{CATT7}MSC group are specific to MIF licensed MSCs.
These data show that MIF licensing can restore MSC immunosuppressive function at doses
that would normally be ineffective.

242 Blocking COX-2 abrogates therapeutic efficacy of MIF licensed BM-MSCs

COX-2 is the rate-limiting enzyme involved in the synthesis of arachidonic acid to PGE2, a 243 key mediator in the immunomodulatory effects of MSCs.⁷⁸ To assess the role of COX-2 on 244 MIF licensed MSCs we inhibited COX-2 with indomethacin. MSCs were treated with 245 246 indomethacin (10 µM) for 30 min. Following the 30 min pre-treatment, cells were incubated 247 with CATT₇ CM for 24 hours. To further validate the involvement of MIF in the improvement 248 of MSC efficacy, MSCs were exposed to an anti-CD74 neutralising antibody (10 µg/ml) or IgG1 isotype control (10 µg/ml) for 30 min. MSCs were then incubated with CATT₇ CM for 249 24 hours (Figure 8A). Analysis of the BAL fluid cell counts showed that pre-treating MSCs 250 with indomethacin before administration significantly reduces ^{CATT7}MSCs ability to suppress 251 252 immune cell infiltration in the BAL fluid of CATT₇ mice challenged with HDM (Figure 8B, 253 8C). Additionally, the analysis of the Th2 cytokines in the BALF showed a marked increase in 254 IL-4 (Figure 8D) and a significant increase in the levels of IL-13 (Figure 8E) in the indomethacin group compared to the control MIF licensed MSC group. Taken together, these 255 256 results show that COX-2 is an important mediator in the enhancement of therapeutic efficacy 257 associated with MIF licensing. Furthermore, blocking of CD74 abrogates MIF licensed BM-258 MSC suppression of eosinophil infiltration and type 2 cytokines in the BAL (Figure 8). These 259 data indicate that MIF enhances MSCs' immunomodulatory capacity mainly through CD74 260 signalling to upregulate COX-2 production.

261

262 Discussion

263 Our main results advance the field of MSC-based therapeutics for asthma by demonstrating 264 that (i) MSC treatment is highly effective in ameliorating airway inflammation; (ii) their

265 therapeutic potential can be enhanced by MSC-MIF licensing as demonstrated in high MIF expressing CATT₇ mice; and finally (iii) that the mechanism of MIF-licensing is dependent 266 267 upon MSC-CD74 expression levels that drive COX-2 expression in MSCs. Our data aligns 268 with the literature demonstrating the ability of human MSCs to ameliorate eosinophil infiltration by reducing the levels of Th2 cytokines.^{8,10-12,25,79} In addition to reducing 269 inflammation, MSCs also alleviate features of airway remodelling in the CATT7 mice. 270 271 Interestingly while MSCs were effective at reducing the severity of goblet cell hyperplasia and 272 subepithelial fibrosis in all groups, we did not observe the same changes in type 2 inflammatory 273 markers in the BAL of WT and low MIF expressing CATT₅ mice suggesting that high levels 274 of MIF may be responsible for improving MSC efficacy. The reduced efficacy of MSCs in the 275 WT and CATT₅ mice is likely attributed to lack of inflammation present associated with a bias 276 towards Th1 immunity in C57BL/6 mice compared to the more Th2 bias in BALB/c mice, influencing the level of Th2 response in our HDM challenge model.⁸⁰ There have been several 277 278 instances where researchers also observed poor responses to MSC treatment of allergic airway 279 inflammation in C57BL/6 mice.^{7,23} More recently, Castro et al. report the requirement of at least 2 doses of human AD-MSCs to reverse airway remodelling and alleviate inflammation in 280 HDM challenged C57BL/6 mice.²¹ 281

We show that a single human MSC dose is capable of significantly decreasing airway 282 283 remodelling in CATT₇ mice. This suggests that high levels of MIF may facilitate activation of 284 MSCs improving their therapeutic efficacy and leading to reversal of airway remodelling. The 285 literature surrounding MSC's effect on airway remodelling is conflicting, however, the majority of the current literature demonstrates that MSCs can attenuate airway 286 remodelling.^{8,10,14,15,17,18,28} Others report a deficit in MSCs capacity to ameliorate goblet cell 287 hyperplasia^{14,19,23} or subepithelial collagen deposition.^{9,28} Reasons for these discrepancies 288 include source of MSCs,^{8,23,81} genotypic mouse model differences, severity of mouse models, 289 time of infusion, MSC fitness, dosing, and route of administration.²⁷ 290

Our previous studies have demonstrated that pro-inflammatory cytokine licensing of MSCs or MSC-like cells; multipotent adult progenitor cells (MAPCs) enhances their retention in inflammatory conditions and correlates with enhanced therapeutic efficacy.^{39,82} We detected significantly higher numbers of MSCs in the lungs of HDM challenged CATT₇ mice compared to CATT₅ or littermate controls 24 hours following administration. It is suggested that shortterm effects of MSCs are mediated by their diverse secretome and the longer-term effects of

MSC therapy are a result of direct interaction with other cell types.⁸³ Increased longevity at the site of injury allows MSCs a longer period to secrete soluble factors and interact with cells in the inflammatory microenvironment. MSC retention in the CATT₇ HDM challenged mice is an important observation and future work will determine if enhanced retention is also involved in the enhanced MSC efficacy observed.

302 Taken together these data suggest that MSCs are more efficacious in the high MIF environment 303 of CATT₇ mice. By investigating the effects of different concentrations of a human cytokine 304 on the efficacy of human MSCs in a model of allergic asthma using a clinically relevant 305 allergen, we have identified a specific disease microenvironment which supports and enhances 306 MSC efficacy. The use of our humanised model aims to provide a more accurate depiction of 307 how human MSCs would interact in subsets of patients compared to conventional murine 308 models. Of course, despite exploring the effect of a human cytokine on human MSCs there are 309 still limitations as we are unable to fully mimic clinical severe allergic asthma and the use of 310 transgenic MIF mice on a C57BL/6 background meant that control WT mice do not develop a 311 high level of type 2 inflammation. However, these results may have implications in tailoring 312 MSC treatment in cases of severe asthma. Our results have demonstrated that MSCs are less efficacious in low MIF environments. Patients with 5/5 haplotypes tend to have lower levels 313 314 of circulating MIF⁸⁴ and therefore may not respond as well to MSC treatment. Whereas patients with 6/6, 7/7 or 8/8 haplotypes are more likely to have high levels of circulating MIF^{52,85,86} 315 316 which may lead to greater MSC activation and enhanced therapeutic efficacy.

317 Following on from the discovery that MSC administration into CATT₇ mice led to improved 318 MSC efficacy, we investigated strategies to use the high MIF microenvironments to potentiate 319 the effects of MSCs. Past work in our lab has focussed on different licensing strategies of MSCs 320 to enhance MSC efficacy. Previously, we have demonstrated how IFN γ licensing can improve 321 MSC efficacy in a humanised model of acute GvHD and how endogenous factors such as peroxisome proliferator-activated receptor (PPAR)δ ligands or treatments like cyclosporine A 322 can influence this.^{39,40} Other studies have shown how licensing with pharmacological agents 323 or endogenous factors can further enhance the effects of MSC therapy in preclinical models of 324 asthma.18,19,87 325

326 One of the main criticisms of preclinical research is the use of doses which far exceed what 327 would be reasonable in the clinic. Analysis of clinical trials using i.v. injection of MSCs reveals 328 that the minimal effective dose used ranges from 1-2 million cells/kg.⁸⁸ Studies which have

investigated the i.v. administration of MSCs in preclinical models of allergic asthma administer doses which equate to 4-40 million cells/kg with the majority at the higher end of the scale.^{8,10,12,16,20-22,25,26,28,79,89-92} The efficacy observed with MIF licensed MSCs using $5x10^4$ cells per mouse results in an effective dose of 2 million cells/kg. This shows that through MIF licensing we are able to restore MSC efficacy at a dose akin to what is used in clinical trials.

334 We then sought to elucidate the mechanisms involved. Given our use of human MSCs in a 335 mouse host, the interspecies ligand/receptor non-functionality can raise questions about how 336 human MSCs might mediate their effects in a mouse host. We and others have shown that indeed human MSCs can mediate protective effects in mouse hosts.^{8,10-12,14-17,21,24,25,33,93} Four 337 338 studies have tracked human MSC biodistribution following i.v. administration in patients with COPD,⁹⁴ liver cirrhosis,⁹⁵ haemophilia A⁹⁶ or breast cancer.⁹⁷ No MSCs were detected in blood 339 at 1hr post infusion. MSCs were distributed mainly in the lung⁹⁵ or lungs and liver⁹⁶ at 48 hr 340 post i.v. infusion with the signal decreasing thereafter. As such, these studies suggest that 341 biodistribution of human MSCs following i.v. administration in humans aligns with the studies 342 investigating MSC biodistribution in mouse models.⁹⁸ 343

344 In terms of mechanism, MIF mediated signal transduction is primarily initiated by binding to MIF's classical receptor CD74.99 We showed that blocking CD74 on the surface of MSCs 345 346 ultimately abolished their immunosuppressive abilities. These findings not only reaffirmed that 347 the licensing with CATT₇ CM was MIF mediated but it also showed that these effects were 348 dependent on binding to CD74. MIF signal transduction through CD74 binding has been shown 349 to initiate a range of signalling pathways which induce cell proliferation, resistance to apoptosis, and the promotion of repair.¹⁰⁰⁻¹⁰⁴ Furthermore, MIF binding to CD74 has been 350 351 shown to activate cytosolic phospholipase A2 (cPLA₂). Moreover, cPLA₂ activation results in the mobilisation of arachidonic acid from membrane phospholipids which is a precursor to the 352 synthesis of prostaglandins.¹⁰⁵ Interestingly MIF can upregulate COX-2 expression, a rate 353 limiting step in the synthesis of prostaglandins such as PGE2,^{71,72,106} however, MIF has been 354 shown to have no effect on the expression of COX-1.71 355

The COX-2/PGE2 pathway has been extensively documented as being one of the key mediators driving MSC immunomodulation.^{39,68,107,108} Our data shows that MIF stimulation enhances the expression of COX-2 but not TSG-6 or PTGS2 in untreated and TNFα licensed MSCs. We hypothesised that the COX-2/PGE2 pathway could be involved in the restoration of MSCs immunomodulatory capacity following CATT₇ licensing. To investigate, we pre-treated MSCs

with indomethacin prior to licensing. Indomethacin is a potent non-selective inhibitor of COX-1 and COX-2.¹⁰⁹ We showed that blocking COX-2 abrogated therapeutic efficacy of CATT₇ licensed MSCs. Interestingly we observed that blocking of COX-2 via indomethacin had more pronounced effect than blocking CD74. COX-2 is constitutively expressed in human MSCs, therefore, inhibition with indomethacin also blocks basal COX-2 expression which will contribute to the effects observed.

These data show that MIF licensing can improve MSC therapeutic efficacy through the 367 upregulation of COX-2 which likely drives PGE2 production. Our data agrees with several 368 369 studies in the literature which also reveal the ability of MIF to improve MSC efficacy in vivo.¹¹⁰⁻¹¹² The Zhang group demonstrated the ability of MIF to improve MSC therapeutic 370 371 efficacy by transducing MSCs with a lentiviral vector containing *Mif* cDNA thus promoting MIF overexpression.¹¹⁰⁻¹¹² Furthermore, Zhang et al. demonstrated the ability of MIF to 372 upregulate COX-2 expression and promote PGE2 production in astrocytes.¹¹³ Here, we further 373 374 demonstrate the effects of ex vivo MIF licensing on MSC therapeutic efficacy by showing binding to CD74 and increased COX-2 expression enhances MSCs immunomodulatory 375 376 abilities.

The knowledge gained from this study can be used to further optimise MSCs as a therapy and provide a basis for future studies regarding the effects of MSCs on the immune response in high MIF environments such as in asthma patients exhibiting the CATT₇ polymorphism.

380

381 Materials and Methods

382 Ethical Approval

All procedures involving the use of animals or human materials were carried out by licensed personnel. Ethical approval for all work was granted by the biological research ethics committee of Maynooth University (BRESC-2018-013). Project authorization was received from the scientific animal protection unit of the health products regulatory agency (HPRA) under AE19124/P022 whereby the terms of the animal experiments within this project were outlined and adhered to in accordance with the ARRIVE criteria.

389 Human bone marrow derived MSC culture

Three different human bone marrow-derived MSC (BM-MSC) donors were obtained from 390 391 RoosterBio Inc. (Frederick, MD, USA). MSCs were first expanded in RoosterBio proprietary 392 expansion medium (RoosterBasal and RoosterBooster) for the first two passages according to 393 manufacturer's instructions. Following this MSCs were cultured and maintained in Dulbecco's 394 Modified Eagles Media Low Glucose (DMEM, Sigma-Aldrich, Arklow, Wicklow, Ireland) 395 supplemented with 10% (v/v) foetal bovine serum (FBS) (BioSera, Cholet, France) and 1% (v/v) Penicillin/Streptomycin (Sigma-Aldrich). Human MSCs were seeded at 1x10⁶ cells per 396 397 T175 flask and cultured at 37 °c in 5% CO₂. Media was replenished every 2-3 days, and cells 398 passaged once they achieved 80% confluency. All experiments were carried out between 399 passages 2-5.

400 MSC Characterisation

401 Three different human bone marrow-derived MSC (BM-MSC) donors (identified as 001-177, 402 003-307 and 003-310) from RoosterBio Inc. (Frederick, MD, USA) were characterised by 403 analysing the expression of cell surface markers. All MSCs donors were found to be negative 404 for CD34 (FITC), CD45 (APC) and HLA-DR (PE) and positive for CD73 (APC), CD90 (FITC) 405 and CD105 (PE) (BD Pharmingen, San Diego, CA, USA) by the Attune Nxt Flow Cytometer 406 (Figure S3).

407 Animal Strains

Two C57BL/6 mouse strains expressing the human high- or low-expression MIF alleles 408 (MIF^{CATT7} [(C57BL/6NTac-Miftm3884.1(MIF)Tac-Tg(CAG-Flpe)2Arte] and MIF^{CATT5} 409 410 [C57BL/6NTac-Miftm3883.1(MIF)Tac-Tg(CAG-Flpe)2Arte] mice) were created using vector-based recombinant replacement of murine Mif by Taconic Biosciences (Rensselaer, 411 412 NY) (Fig. 1). Validation of the expression of human and not murine MIF mRNA was verified 413 by qPCR, and -794 CATT-length dependent stimulated MIF production was confirmed in vivo.⁶¹ Littermate wildtype (WT) and MIF^{-,-} (MIF KO)¹¹⁴ (a kind donation from R. Bucala, 414 415 Yale School of Medicine, Yale University, New Haven, CT, USA) mice were used as controls.

416 HDM-induced Airway Inflammation Model and Therapeutic Protocol

Both male and female MIF^{CATT7}, MIF^{CATT5} or WT mice aged 6-12 weeks were challenged with 417 25 µg HDM extract (Dermatophagoides pteronyssinus, Greer Laboratories, Lenoir, NC, USA) 418 419 in 25 µl phosphate buffered saline (PBS) intranasally (i.n.) 3 days weekly for 3 weeks under 420 light isoflurane anaesthesia. Control mice were challenged with 25 µl PBS under the same 421 conditions. On day 14, after HDM challenge, mice received an intravenous (i.v.) injection of 1×10^6 MSCs in 300 µl into the tail vein.¹¹⁵ For the dose curve 1×10^6 , 5×10^5 , 1×10^5 , and 5×10^4 422 were administered i.v. into HDM challenged CATT₇ mice on day 14. $5x10^4$ was selected as the 423 424 dose at which MSCs lose efficacy.

425 Licensing of MSCs with Endogenous Human MIF

Supernatants containing endogenous human MIF were generated from bone marrow- derived 426 427 macrophages (BMDMs) of C57BL/6 mouse strains expressing the high-expressing MIF allele 428 (CATT₇). CATT₇ mice were challenged with HDM in 25 µl phosphate buffered saline (PBS) 429 intranasally 3 days weekly for 3 weeks under light isoflurane anaesthesia. 4 hr post final challenge, femurs and tibias were flushed with warm Roswell Park Memorial Institute (RPMI) 430 1640 medium GlutaMAXTM (Gibco, Paisley, UK) supplemented with 10% (v/v) heat 431 432 inactivated foetal bovine serum (FBS) (BioSera) and 1% (v/v) Penicillin/Streptomycin (Sigma-433 Aldrich). Cells were collected and seeded into T175 flasks in cRPMI supplemented with 10% 434 L929 conditioned medium. L929 cell line produces high amounts of macrophage colony 435 stimulating factor (M-CSF) and other proteins stimulating macrophage differentiation. After 436 96 hours, supernatants were collected, sterile filtered (0.22 μ M pore size) and stored at -20°C. 437 The conditioned media generated in this manner will be referred to as CATT₇ CM. 438 Additionally, KO CM was generated from MIF KO mice as a control. Licensing MSCs was performed by removing existing media, washing with PBS, and incubating cells with CATT₇ 439 CM (^{CATT7}MSC) or MIF KO CM (^{KO}MSC) for 24 hr. To account for variability of human MIF 440 441 levels between CATT₇ mice and to ensure WT mice did not produce human MIF, CATT₇, 442 CATT₅ and WT supernatants were measured by human MIF ELISA (R&D Systems, MN, USA) as described previously (Figure S2). 62 5x10⁴ licensed MSCs were administered i.v. into 443 444 HDM challenged CATT₇ mice on day 14. Where indicated, MSCs were treated with COX-2 445 inhibitor indomethacin (10 µM) for 30 min. Following pre-treatment, MSCs were licensed with 446 CATT₇ CM for 24 hr as described above. Moreover, mouse anti-CD74 neutralising antibody

and isotype control were added to the assay. MSCs were pre-treated with anti-CD74 neutralising antibody (clone LN2) (10 μ g/ml) or IgG1 κ isotype control (clone T8E5) (10 μ g/ml) for 30 min. MSCs were then licensed with CATT₇ CM for 24 hr before administration.

450 Collection of Bronchoalveolar Lavage (BAL) Fluid

On day 18, 4 hr post final challenge, mice were sacrificed for cell and cytokine analysis of the 451 452 BAL fluid. BAL fluid was obtained through 3 gentle aspirations of PBS. After centrifugation, protease inhibitor cocktail (Roche Diagnostics, Mannheim, Germany) was added to the 453 454 supernatants before Th2 cytokine analysis. Total numbers of viable BAL cells were counted 455 using ethidium bromide/acridine orange staining on a haemocytometer then pelleted onto 456 microscope slides by cytocentrifugation. Slides were stained with Kwik Diff kit stain (Richard-457 Allan Scientific, Kalamazoo, MI, USA) and coverslips were mounted using DPX mounting 458 medium (Sigma-Aldrich). Differential cells counts were derived by counting a minimum of 459 300 leukocytes on randomly selected fields under a light microscope at 20X magnification.

460 Enzyme-linked immunosorbent assay (ELISA)

Levels of mIL-4 (Biolegend, San Diego, CA, USA) and mIL-13 (eBioscience, San Diego, CA,
USA) were determined using commercial ELISA kits, according to manufacturer's
instructions.

464 Lung Histology

On day 21, mice were sacrificed for histological analysis. Lungs were removed and fixed in 465 466 10% neutral buffered formalin, paraffin embedded and 5 µm slices were mounted onto slides 467 for histological analysis. Lung tissue was stained with haematoxylin and eosin (H&E), periodic 468 acid-Schiff (PAS) or Masson's Trichrome to analyse immune cell infiltration, goblet cell hyperplasia or subepithelial collagen deposition respectively. H&E analysis was carried out as 469 previously described.¹¹⁶ Goblet cell hyperplasia was determined by the % of PAS positive cells 470 471 in airways relative to airway diameter. Subepithelial collagen deposition was calculated by 472 analysing the % of positive staining using the trainable Weka segmentation plugin on Fiji open-473 source software.

474 Cryo-Imaging

1x10⁶ MSCs were labelled with the Qtracker® 625 labelling kit (Invitrogen, Paisley, UK) 475 476 according to manufacturer's instructions before being administered i.v. on day 14. On day 15, 477 mice were humanely euthanised, and the lungs were embedded in mounting medium for 478 cryotomy (O.C.T compound, VWR Chemicals, Leuven, Switzerland), frozen in liquid nitrogen 479 and stored at -80°c. Lungs were sectioned into 40 µm slices and imaged with the automated CryoViz[™] imaging system (BioInvision Inc., Cleveland, OH, USA). Images were then 480 481 processed to generate 3D images using CryoViz[™] processing, and the number of detected cells was quantified using cell detection software (BioInvision).³⁹ 482

483 Analysis of Gene Expression

Total RNA was extracted using TRIzol (Ambion Life Sciences, Cambridgeshire, UK) 484 485 according to manufacturer's instructions. RNA concentrations were measured using a 486 spectrophotometer (Nanodrop 2000, ThermoScientific, Wilmington DE, USA) and were 487 equalised to 100ng/µl before cDNA synthesis. cDNA synthesis was performed using 488 manufacturer's instructions (Quantabio, MA, USA). Real Time-Polymerase Chain Reaction 489 (RT-PCR) was carried out using PerfeCta SYBR Green FastMix (Quantbio, MA, USA). 490 Expression of human COX-2, PTGES, IDO, ICAM-1, HGF, TSG-6 and PTGS2 (primer 491 sequence information is contained in Table S1) was quantified in relation to the housekeeper 492 gene HPRT using the ΔCT method. The fold change in the relative gene expression was determined by calculating the $2^{-\Delta\Delta CT}$ values. 493

494 MSC Expansion Assay

495 1.4×10^{3} MSCs were seeded out into T25 flasks in cDMEM or 50:50 cDMEM and WT CM or 496 CATT₇ CM for 72 hr. Cells were trypsinised and stained with ethidium bromide/ acridine 497 orange and counted on a haemocytometer. MIF inhibitor SCD-19 (100 μ M) was used to 498 determine MIF specificity. In such cases, conditioned media was pre-incubated with SCD-19 499 1 hr before the expansion assay.

500 Intracellular Staining of COX-2 and IDO

501 MSCs were seeded at 1×10^5 cells per well in 6 well plates. MSCs were stimulated with 502 recombinant human IFN γ at low (5ng/ml) or high (40ng/ml) concentrations, TNF α (5-10ng/ml)

- 503 (PeproTech, London, UK), recombinant human MIF (1ng/ml) (provided by Rick Bucala, Yale)
- 504 or endogenous MIF (CATT₇ CM) for 24 hr. Cells were prepared for intracellular staining using
- 505 the Intracellular FoxP3 kit as per manufacturer's instructions. Cells were stained with COX-2
- 506 (PE) or IDO (APC) (BD Pharmingen, San Diego, CA, USA) for 45 min. Cells were then
- 507 washed in flow cytometry staining buffer and acquired using the Attune Nxt Flow Cytometer.

508 Surface Staining of CD74

509 MSCs were seeded at 1×10^5 cells per well in 6 well plates. MSCs were stimulated with 510 recombinant human IFN γ (5ng/ml) (PeproTech, London, UK), recombinant human MIF (1, 10

- 511 or 100ng/ml) or endogenous MIF (CATT₇ CM) for 24 hr. Cells were stained with CD74 (PE)
- 512 (BD Pharmingen, San Diego, CA, USA) for 45 min. Cells were then washed in flow cytometry
- 513 staining buffer and acquired using the Attune Nxt Flow Cytometer.

514 T Cell Suppression assay

Human PBMCs were isolated from buffy packs (Irish Blood Transfusion Service), by Ficoll 515 density gradient centrifugation. 5×10^4 Carboxyfluorescein succinimidyl ester (CFSE) labelled 516 PBMC were co-cultured (ThermoFisher Scientific, Eugene, OR, USA) with BM-MSC in a 517 1:20 ratio $(2.5 \times 10^3 \text{ cells/well})$. 24 hr prior to co-culture, BM-MSCs were incubated with 518 CATT₇ CM or CATT₇ CM + SCD-19 (100 µM). After 24 hr, BM-MSCs were washed with 519 520 PBS before adding the PBMCs. Activation and expansion of human T cells was carried out using ImmunoCultTM human CD3/CD28 T cell activator antibody mix (STEMCELL 521 Technologies, Cambridge, UK). After 4 days, PBMCs were harvested and the frequency (%) 522 523 and number of proliferating CD3⁺ T cells were analysed by flow cytometry (Attune Nxt Flow 524 Cytometer).

525 Statistical Analysis

Mice were randomised. Observers assessing end-points were blinded to group assignment.
Data for individual animals and independent experiments are presented as individual symbols.
All data are presented as mean ± SEM. Results of two or more groups were compared by oneway analysis of variance (ANOVA) followed by the *post-hoc* Tukey's multiple comparison
test. GraphPad Prism (GraphPad Software Inc, San Diego, CA, USA) was used for all statistical
analyses.

532 Data Availability Statement

The data that support the findings of this study are available on request from the correspondingauthor.

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542 Author's Contributions

543 IJH performed research, data analysis, study design and wrote the manuscript. HD performed 544 research, data analysis, study design and wrote the manuscript. CT & TS performed research 545 and data analysis; DJW, SRE & CCDS contributed to study design and data analysis. SCD & 546 MEA provided reagents, contributed to study design and data analysis. KE designed and 547 supervised the study and wrote the manuscript. All authors approved the final manuscript.

548 **Declaration of Interest**

549 The authors declare no conflict of interest.

550 Keywords

- Mesenchymal Stromal Cells, House Dust Mite, Allergic Asthma, Macrophage Migration
 Inhibitory Factor, Cyclooxygenase

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957

958 List of Figure Captions

959 Figure 1. Human BM-MSCs significantly reduce goblet cell metaplasia and collagen 960 deposition in CATT₇ mice challenged with HDM. A PBS and HDM groups received PBS or 961 HDM i.n. 3 times a week for 3 consecutive weeks. 1x10⁶ human BM-MSCs were administered 962 i.v. to the HDM+MSC groups on day 14. Mice were sacrificed on day 21 (Schematic created 963 with BioRender.com). B Representative images of lung tissue from WT, CATT₅ and CATT₇ mice stained with Periodic acid Schiff (PAS) at 20X magnification, scale bar = $20 \,\mu$ m. Arrows 964 965 show examples of mucin-containing goblet cells. C Goblet cell hyperplasia was investigated 966 through the quantitation of PAS positive cells. D Representative images of lung tissue stained 967 with Masson's trichome at 4X magnification, scale bar = $200 \mu m$. E Quantitation of % subepithelial collagen. Data are presented as mean \pm SEM; N=6 per group. Human BM-MSC 968 969 donors 001-177 and 003-310 were used (RoosterBio Inc., Frederick, MD, USA). Statistical analysis was carried out using one-way ANOVA followed by the post-hoc Tukey's multiple 970 comparison test where *p<0.05, ***p<0.001, ****p<0.0001, ns: non significant. 971

972

Figure 2. Human BM-MSCs significantly reduce levels of Th2 cytokines in the BAL fluid 973 974 of CATT₇ mice challenged with HDM. PBS and HDM groups received PBS or HDM i.n. 3 times a week for 3 consecutive weeks. 1x10⁶ human BM-MSCs were administered i.v. to the 975 976 HDM+MSC groups on day 14. BAL was performed 4 hr post final HDM challenge on day 18. 977 A Total cell count recovered from the BAL. B BAL fluid eosinophil count determined by 978 differential staining of cytospins. Cytokine levels of C IL-4 and D IL-13 in the BAL fluid 979 determined by ELISA. White bar: PBS; Grey bar: HDM; Blue bar: HDM+MSC. Data are 980 presented as mean ± SEM; N=5-6 per group. Human BM-MSC donors 001-177 and 003-310 981 were used (RoosterBio Inc., Frederick, MD, USA). Statistical analysis was carried out using 982 one-way ANOVA. followed by the *post-hoc* Tukey's multiple comparison test where *p<0.05, **p<0.01, ***p<0.001, ****p<0.0001, ns: non significant. 983

984

985 Figure 3. High levels of hMIF significantly enhance BM-MSC retention in a HDM model

986 of allergic asthma. HDM was administered i.n. 3 times a week for 2 weeks. On day 14, $1x10^6$

987 Qtracker 625- labelled hMSCs were administered i.v. in WT, CATT₅ or CATT₇ mice. 24 hr

later the lungs were harvested, embedded in OCT compound and frozen at -80. Tissue blocks
 were sectioned and imaged using the CryoVizTM (Bioinvision, Cleveland, OH, USA) imaging

990 system. 3D images show representative lung images from **A** WT, **B** CATT₅ and **C** CATT₇ mice 991 with detected MSCs shown in yellow. **D** Total number of MSCs detected in the lung and **E** 992 number of clusters were quantified using CryoVizTM Quantification software. Data are 993 presented as mean \pm SEM; N=3 per group. Human BM-MSC donor 001-177 was used 994 (RoosterBio Inc., Frederick, MD, USA). Statistical analysis was carried out using one-way 995 ANOVA followed by the *post-hoc* Tukey's multiple comparison test where **p<0.01.

996 Figure 4. Influence of rhMIF licensing on MSC expression of immunomodulatory factors

997 *in vitro*. **A-E** Gene expression of IDO, COX-2, PTGES, ICAM-1 and HGF by hBM-MSCs 998 after stimulation with recombinant human MIF (1ng/ml), human TNF α or human IFN γ for 999 24hr. Data are presented as mean ± SEM and are representative of 3 independent experiments. 1000 Human BM-MSC donors 001-177, 003-310 and 003-307 were used (RoosterBio Inc., 1001 Frederick, MD, USA). Statistical analysis was carried out using one-way ANOVA where 1002 *p<0.05, **p<0.01, ****p<0.0001, ns: non significant.

1003

Figure 5. CATT₇ MIF licensing enhances MSC expansion and immunosuppressive function *in vitro*.

1006 A Schematic (created using Biorender.com) depicting the generation of CATT₇ MIF CM and 1007 experimental design. B-E Percentage or mean fluorescence intensity (MFI) of IDO or COX-2 1008 expression in human BM-MSCs measured by flow cytometry after cells were stimulated with 1009 CATT₇ MIF CM, human TNF α or human IFN γ for 24hr. F Percentage expression and 1010 representative histogram plots of CD74 surface expression on human MSCs measured by flow 1011 cytometry after cells were stimulated with CATT₇ MIF CM and human IFN_γ for 24hr. G-H 1012 Relative gene expression of TSG-6 and PTGS2 by hBM-MSCs after cells were stimulated with 1013 endogenous human MIF (CATT₇ CM) and human TNF α for 6hr. I-J Licensing of MSCs with 1014 supernatants generated from BMDMs from CATT₇ HDM challenged mice enhances MSC 1015 suppression of (I) frequency (%) and (J) absolute number of $CD3^+$ T cells proliferating. Blockade of MIF using SCD-19 (100 µM) in the BMDM supernatants 1 hr before addition to 1016 1017 MSCs abrogates the enhanced effect of MIF on MSC suppression of T cell proliferation (I-J). 1018 K Licensing of MSCs with CATT₇ MIF CM enhances MSC expansion in vitro. Addition of 1019 MIF inhibitor SCD-19 (100 µM) to CATT₇ MIF CM 1 hr before MSC licensing prevents MIF 1020 enhanced MSC expansion. Data are presented as mean \pm SEM and are representative of 3

independent experiments. Human BM-MSC donors 001-177, 003-310 and 003-307 were used
(RoosterBio Inc., Frederick, MD, USA). Statistical analysis was carried out using a one-way
ANOVA or unpaired t test where *p<0.05, **p<0.01, ***p<0.001, ****p<0.0001, ns: non
significant.

1025

1026 Figure 6. Titration of BM-MSC doses in CATT₇ mice challenged with HDM. A To 1027 determine the point where MSCs lose efficacy in CATT₇ mice, a range of doses were 1028 administered on day 14. BAL was performed 4 hr post final HDM challenge on day 18 1029 (Schematic created with BioRender.com). B Total cell count recovered from the BAL. C 1030 Number of eosinophils obtained from the BAL fluid. Cytokine levels of **D** IL-4 and **E** IL-13 in 1031 the BAL fluid determined by ELISA. Data are presented as mean \pm SEM; N=2-3 per group. 1032 Human BM-MSC donors 001-177 and 003-310 were used (RoosterBio Inc., Frederick, MD, 1033 USA). Statistical analysis was carried out using one-way ANOVA followed by the post-hoc 1034 Tukey's multiple comparison test where p<0.05.

1035

Figure 7. MIF licensing restores MSC efficacy at low doses in CATT₇ mice. A 5x10⁴ MSCs 1036 were administered to HDM challenged CATT₇ mice on day 14. ^{CATT7}MSCs were licensed with 1037 CATT₇ BMDM supernatant for 24 hr prior to i.v. administration. The control group ^{KO}MSCs 1038 1039 were generated by licensing MSCs with BMDM supernatant from MIF KO mice 24 hr prior to i.v. administration. BAL was performed 4 hr post final HDM challenge on day 18 (Schematic 1040 created with BioRender.com). B Total number of cells in the BAL were determined and 1041 1042 differential cell counts were performed on the collected cells to determine the numbers of C 1043 eosinophils. Cytokine levels of **D** IL-4 and **E** IL-13 in the BAL fluid determined by ELISA. 1044 Data are presented as mean \pm SEM; N=5-6 per group. Human BM-MSC donors 001-177 and 1045 003-310 were used (RoosterBio Inc., Frederick, MD, USA). Statistical analysis was carried out 1046 using one-way ANOVA followed by the post-hoc Tukey's multiple comparison test where 1047 *p<0.05.

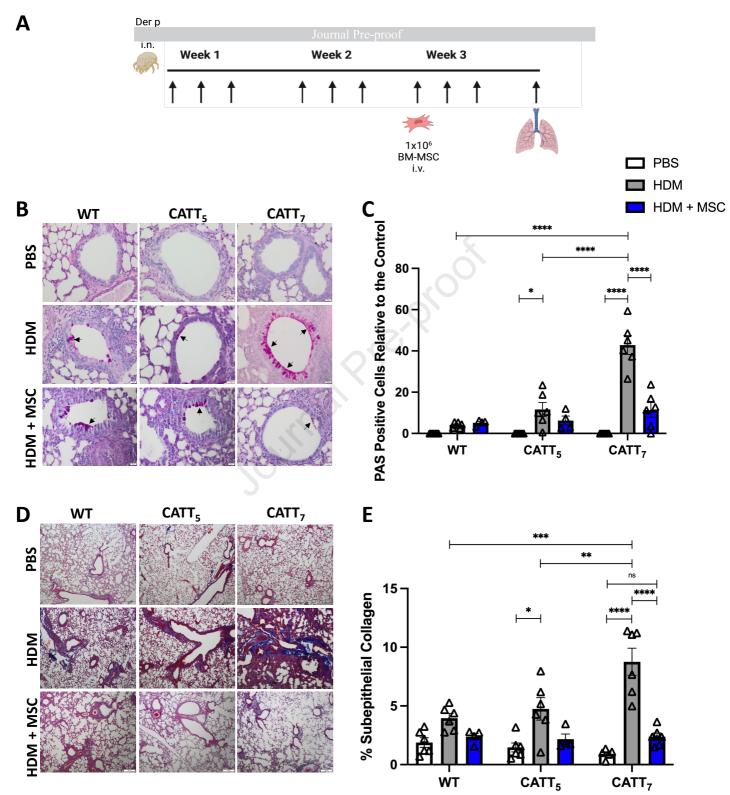
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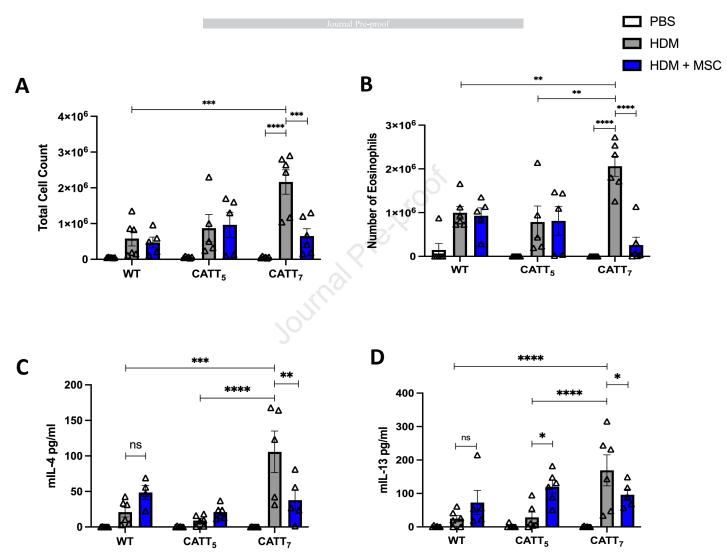
Figure 8. MIF-Licensed MSCs mediate their protective effects in HDM-induced allergic
 airway inflammation in a CD74 and COX-2 dependent manner in CATT₇ mice. A 5x10⁴

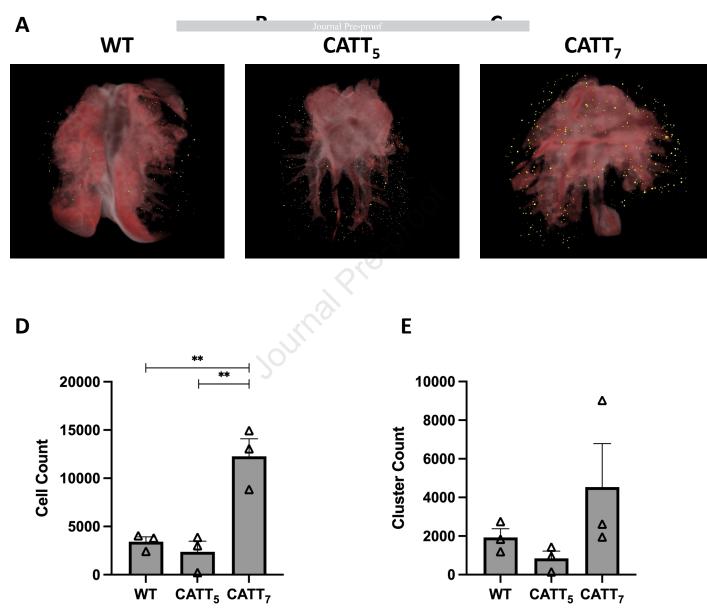
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1051 MSCs were exposed to the COX-2 inhibitor indomethacin, an anti-CD74 neutralising antibody 1052 or an isotype control antibody for 24 hr in vitro. All MSCs were licensed with CATT7 BMDM 1053 supernatant for 24 hr prior to i.v. administration. BAL was performed 4 hr post final HDM 1054 challenge on day 18 (Schematic created with BioRender.com). B Total number of cells in the 1055 BAL were determined and differential cell counts were performed on the collected cells to 1056 determine the numbers of C eosinophils. Cytokine levels of D IL-4 and E IL-13 in the BAL 1057 fluid determined by ELISA. Data are presented as mean \pm SEM; N=5-6 per group. Human 1058 BM-MSC donors 001-177 and 003-310 were used (RoosterBio Inc., Frederick, MD, USA). 1059 Statistical analysis was carried out using one-way ANOVA followed by the *post-hoc* Tukey's 1060 multiple comparison test where *p<0.

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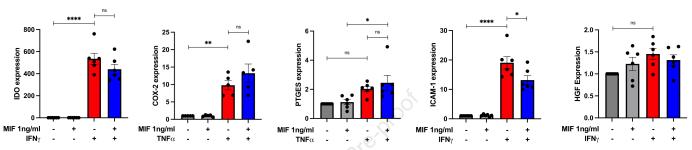






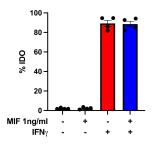
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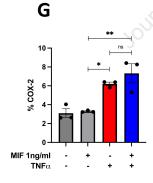


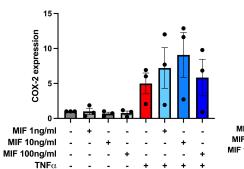
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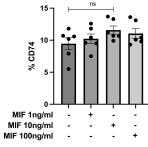


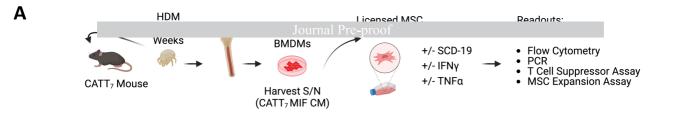
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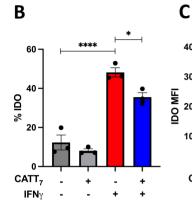


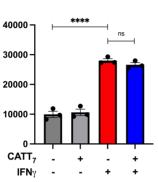


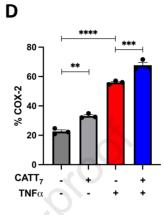
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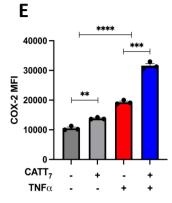


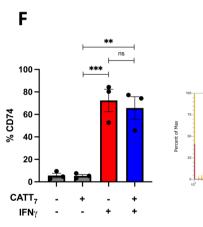


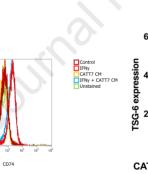


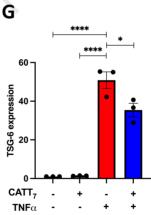












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