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Cortisol facilitates immune escape of human acute myeloid leukaemia cells by inducing

2	latrophilin 1 expression
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23 Progression of acute myeloid leukaemia (AML) - the most severe blood/bone marrow cancer - is determined by the ability of malignant cells to escape host immune surveillance. 24 However, the systemic regulation mechanisms underlying this phenomenon remain largely 25 26 unknown. In this study we have discovered a fundamental systemic biochemical strategy which allows AML cells to employ physiological body systems to survive and escape 27 immune attack. We found that AML cells use a crucial human adrenal cortex hormone 28 (cortisol) to induce the expression of the neuronal receptor latrophilin 1 (LPHN1) which 29 facilitates exocytosis. This receptor interacts with the blood plasma protein FLRT3 30 31 (fibronectin leucine rich transmembrane protein 3) to cause secretion of the immune suppressor galectin-9 which impairs the anti-cancer activities of cytotoxic lymphoid cells. 32

AML is a cancer of the blood and bone marrow which originates from self-renewing 33 malignant immature myeloid cells and rapidly becomes a systemic, and very often fatal, 34 malignancy¹. AML cells employ body systems to produce factors required for their 35 proliferation/disease progression^{2,3}. This includes employment of stem cell factor (SCF), a 36 37 major hematopoietic growth factor controlling AML progression and thus becoming highly oncogenic^{2,3}. Expression and release of SCF can be triggered by AML cells via cytokines 38 (e.g. interleukin-1 β)². Recent evidence clearly demonstrated that AML cells are also capable 39 of impairing the activities of cytotoxic lymphoid cells (e.g. natural killer (NK) cells and 40 cytotoxic T cells⁴). One of the biochemical mechanisms underlying this phenomenon lies in 41 the ability of AML cells to secrete the protein called galectin-9. This tandem type galectin 42 binds the immune receptor Tim-3 and induces a variety of intracellular and cell-to-cell 43 signalling events leading to the inactivation of NK cells as well as killing of cytotoxic T 44 cells^{4,5}. We recently reported that the process of galectin-9 secretion in AML cells is 45 46 stimulated by the unique G protein-coupled receptor LPHN1, which normally functions in neurons facilitating exocytosis^{4,6}. LPHN1 is also found in haematopoietic stem cells (HSCs) 47

but its expression disappears at early stages of their maturation^{4,7}. However, upon malignant
transformation, AML cells preserve their abilities to express LPHN1 and produce high levels
of galectin-9 and Tim-3, where the latter is involved in trafficking galectin-9 during the
secretion process (HSCs express neither galectin-9 nor Tim-3⁴).

It is currently unknown which molecular mechanisms trigger high levels of LPHN1 expression in primary human AML cells, and in general mechanisms of upregulation of LPHN1 expression at the genomic level remain unclear. It is also unknown whether FLRT3, a natural LPHN1 ligand^{4,8}, is present in human blood plasma and other tissues associated with AML. Unravelling these mechanisms is crucial in order to understand the pathways that control the ability of AML cells to protect themselves against cytotoxic lymphoid cells and thus was the aim of the present study.

59 **Results and Discussion**

In order to investigate the effects of cortisol on LPHN1 transcription, we exposed primary 60 and THP-1 human AML cells, primary human HSCs and primary healthy human leukocytes 61 to 1 µM cortisol for 24 h followed by quantitative real-time PCR analysis of LPHN1 mRNA 62 levels. We found that all cell types, except primary healthy leukocytes, transcribed detectable 63 64 amounts of LPHN1 mRNA and in all these cases levels were significantly upregulated by treatment with cortisol (Figure 1A). In both THP-1 and primary human AML cells LPHN1 65 66 protein levels were also clearly upregulated (Figure 1B and C). In contrast, primary human healthy leukocytes did not express detectable amounts of LPHN1 protein and this was not 67 68 altered by the effects of cortisol (Figure 1D). Comparative analysis of LPHN1 protein expression in primary human AML cells, THP-1 cells and primary human healthy leukocytes 69 70 is shown in Supplementary figure 1.

Cortisol treatments did not upregulate galectin-9 secretion in any of these cell types (Figure 1
B-D), suggesting that LPHN1 needs to be activated by a ligand to induce galectin-9 release.

Analysis of blood plasma levels of cortisol in AML patients vs healthy donors (samples were 73 collected at the same time of the day to avoid the influence of circadian dynamics) 74 demonstrated that its levels were significantly higher in the blood plasma of AML patients 75 compared to healthy donors (Figure 1E). Galectin-9 levels were also substantially higher in 76 AML patients (Figure 1F), which is in line with our previous observations⁴. Furthermore, 77 there was no correlation between cortisol and galectin-9 levels in the blood plasma of healthy 78 donors, while in AML patients there was a clear correlation (Figure 1G), suggesting that 79 80 galectin-9 secretion in this case might be linked to LPHN1 expression.

81 If LPHN1 is expressed on the surface of blood cells, it can also be shed by proteolysis and therefore appear in the plasma. LPHN1 was immunoprecipitated, extracted and subjected to 82 Western blot analysis from blood plasma samples from AML patients using several LPHN1 83 84 antibodies. A clear fragment was detectable at around 67-68 kDa, smaller fragments were 85 detectable as well, but only in AML plasma, while in the blood plasma of healthy donors there was no evidence of the presence of LPHN1 fragments (Figure 1H). These fragments 86 were also detectable by ELISA (Figure 1I, see Materials and Methods for description of the 87 ELISA format). 88

As reported before⁴, we observed that exposure of THP-1 AML cells to 10 nM FLRT3 for 16 h resulted in a significant increase in galectin-9 secretion (Figure 2A). This effect was not detectable in primary healthy human leukocytes (Figure 2A). Importantly, 1 h pre-exposure of THP-1 cells to rabbit polyclonal antibody recognising LPHN1 (clone name RL1⁹) prior to 16 h treatment with 10 nM FLRT3 attenuated FLRT3-induced galectin-9 release, confirming involvement of LPHN1 in this process (Figure 2A). The antibody employed specifically 95 recognised target molecules on the surface of THP-1 cells (Supplementary figure 2). We used 96 mouse neuroblastoma cells NB2A, which do not express LPHN1¹⁰, as a negative control and 97 measured the interaction of the antibody with the cell surface using a Li-Cor on-cell assay as 98 described in the Materials and Methods (please see supplementary information). Exposure of 99 THP-1 cells to 1 μ g/ml RL1 for 16 h did not affect galectin-9 secretion levels (data not 90 shown) suggesting that this antibody lacks a LPHN1 agonistic effect.

101 Interestingly, we found that blood plasma of both healthy donors and AML patients contains approximately equal amounts of secreted FLRT3 (most likely by proteolytic shedding) with a 102 molecular weight of approximately 55 kDa (which corresponds to the molecular weight of 103 FLRT3 shed from the cell surface by proteinases¹¹). Another specific band was seen at 104 around 27-28 kDa which most likely corresponds to a smaller cleavage fragment of the 105 FLRT3 extracellular domain (Figure 2B). The amounts of this smaller fragment were also 106 107 equal in blood plasma of healthy donors and AML patients (Figure 2B). To explore which blood plasma-based ligands can induce galectin-9 secretion in AML cells we cultured THP-1 108 109 cells in RPMI-1640 medium containing antibiotics (as outlined in Materials and Methods see supplementary information) replacing 10 % foetal bovine serum (FBS) with blood plasma 110 from either healthy donors or AML patients. Cells were incubated for 16 h with or without 30 111 min pre-incubation with anti-FLRT3 antibody in order to neutralise FLRT3 activity. 112 Galectin-9 secretion levels were significantly higher in the presence of both sources of human 113 blood plasma compared to FBS (negative control). Anti-FLRT3 antibody attenuated galectin-114 9 secretion (Figure 2C). Binding of LPHN1 and FLRT3 was further confirmed using SRCD 115 spectroscopy. We found that the two proteins interact with each other with high affinity, 116 inducing conformational change in both proteins, as seen from far UV synchrotron radiation 117 circular dichroism (SRCD) spectra (Figure 2D). This is further confirmation of the high 118

affinity interaction of LPHN1 and FLRT3 observed in previous studies⁸ using different
techniques.

Taken together, our results demonstrate, for the first time, that cortisol upregulates LPHN1 121 expression at the transcriptional level, thus stimulating its translation in human AML cells. 122 AML leads to a decreased blood plasma glucose levels⁵, which normally leads to 123 upregulation of secretion of corticotropin-releasing hormone (CTRH) by hypothalamus¹². 124 CTRH induces secretion of adrenocorticotropic hormone (ACTH) by pituitary gland¹². 125 ACTH upregulates cortisol production by the adrenal cortex¹². Cortisol is then employed by 126 AML cells. In healthy human leukocytes cortisol is not capable of inducing LPHN1 127 transcription/translation, possibly because of gene repression. Interaction of AML cell-128 derived LPHN1 with released FLRT3 available in blood plasma facilitates the secretion of 129 galectin-9. The latter protects AML cells against immune attack which could otherwise be 130 131 performed by NK cells as well as cytotoxic T cells (Supplementary figure 3). Importantly, LPHN1 fragments are present in the blood plasma of AML patients but not in healthy donors. 132 These fragments were detectable by both Western blot analysis and ELISA, which indicates 133 the possibility of detection of these fragments for rapid AML diagnosis, although differential 134 verification tests have yet to be performed. Our results suggest a fundamentally novel 135 mechanism used by AML cells in order to progress the disease. They use a common 136 endogenous human hormone (cortisol) to induce LPHN1 expression by employing a widely 137 available ligand (FLRT3, which is always present in blood plasma) in order to escape host 138 immune surveillance. Thus, AML cells employ crucial functional systems of the human body 139 to support their survival and attenuate the anti-cancer activities of cytotoxic lymphoid cells. 140 Our work indicates that galectin-9 and secreted FLRT3 are the most promising targets for 141 anti-AML immune therapy. 142

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145	Materials and Methods are presented in Supplementary Information.						
146							
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150	Source for access to B23 beamline (SM12578).						
151							
152	Conflict of interest						
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154	The Authors have no conflict of interest to declare						
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215 Figure legends.

Figure 1. Cortisol induces LPHN1 expression in human AML cells and haematopoietic 216 stem cells but not in primary healthy human leukocytes. Primary human AML, THP-1 217 and haematopoietic stem cells as well as primary healthy leukocytes were exposed to 1 µM 218 219 cortisol for 24 h followed by analysis of LPHN1 gene transcription by quantitative real-time PCR (A) and Western blot analysis (B – primary AML cells, C – THP-1 cells and D – PHL). 220 For PHL, lysates of LPHN1 overexpressing NB2A cells were used as a positive control. 221 ELISA was used to measure secreted galectin-9 levels. Blood plasma of ten healthy donors 222 and ten AML patients was collected at the same time of the day to ensure comparability of 223 224 cortisol levels. Cortisol (E) and galectin-9 (F) levels were measured by ELISA and correlation between the levels of these two proteins was analysed (G). Soluble LPHN1 225 fragments were immunoprecipitated and detected by Western blot (H) and ELISA (I), as 226 outlined in Materials and Methods. Images are from one experiment representative of four – 227 six which gave similar results. Data represent mean values \pm SEM of six – ten independent 228 experiments. ; *p < 0.05; **p < 0.01; ***p < 0.01 vs. control. 229

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Figure 2. FLRT3 induces galectin-9 secretion in AML cells in a LPHN1-dependent manner. THP-1 cells and PHL were exposed to 10 nM human recombinant FLRT3 for 16 h, followed by detection of secreted galectin-9 by ELISA. In THP-1 cells, the treatment was performed with or without 1 h pre-exposure to 1 µg/ml RL1 anti-LPHN1 polyclonal antibody

235 (A). The levels of released FLRT3 fragments were analysed in the blood plasma of healthy donors and AML patients using Western blot (B). THP-1 cells were exposed for 16 h to 10% 236 blood plasma either from healthy donors or AML patients, with or without pre-treatment with 237 238 FLRT3 neutralising antibody. Levels of secreted galectin-9 were analysed using ELISA. (C). Secondary structure and conformational changes of LPHN1, FLRT3 and the complex of the 239 two proteins were characterised using SRCD spectroscopy as outlined in Materials and 240 Methods (D). Images are shown from one representative experiment of four which gave 241 similar results. Data are shown as mean values \pm SEM from four independent experiments; 242 *p < 0.05; **p < 0.01 vs. control. 243

244

245 **Figure 1**







253 latrophilin 1 expression

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276	Materials and Methods								
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278	Materials								
279	RPMI-1640 medium, foetal bovine serum, supplements as well as basic laboratory chemicals								
280	were purchased from Sigma-Aldrich (Suffolk, UK). Maxisorp™ microtitre plates were								
281	obtained from Nunc (Roskilde, Denmark) and Oxley Hughes Ltd (London, UK). Human								
282	recombinant FLRT3, mouse monoclonal antibody against LPHN1 and rabbit antibody against								
283	FLRT3 were purchased from Santa Cruz Biotechnology (Dallas, Texas, USA). PAL1 and								
284	RL1 rabbit polyclonal antibodies against LPHN1 were described previously ^{1,2} . Rabbit								
285	antibody against native LPHN1 was obtained from Abcam (Cambridge, UK). Goat anti-								

286 mouse and goat anti-rabbit fluorescence dye-labelled antibodies were obtained from LI-COR

287 (Lincoln, Nebraska USA). ELISA-based assay kits for the detection of galectin-9 were purchased from Bio-Techne (R&D Systems, Abingdon, UK). A soluble extracellular 288 fragment of LPHN1, LPH-51, was produced and purified as described before². Briefly, NB2a 289 290 cells stably expressing LPH-51 were cultured in serum-free medium for 48 h. 30 mL of the medium was then incubated overnight at 4°C with 500 µL of anti-V5-antibody agarose 291 (Sigma-Aldrich). After incubation, the column was washed with 10 column volumes of PBS. 292 Bound protein was fractionally eluted with 5 volumes of elution buffer containing 50 mM 293 triethylamine and 150 mM NaCl (pH 12). The eluted protein was immediately neutralized 294 with 150 mM NaCl, 1 M HEPES (pH 7.2)⁹. The fractions were then analysed by Western 295 blotting and combined. The protein was concentrated using Amicon centrifugal ultrafiltration 296 units with a 30,000 molecular weight cut-off (Sigma-Aldrich). All other chemicals purchased 297 298 were of the highest grade of purity commercially available.

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300

301 Cell lines and primary cells

THP-1 human myeloid leukemia monocytes cells were obtained from the European Collection of Cell Cultures (Salisbury, UK). Cells were cultured in RPMI 1640 medium (R8758 – Sigma-Aldrich) with L-glutamine and sodium bicarbonate, liquid, sterile-filtered, suitable for cell culture) supplemented with 10% foetal bovine serum, penicillin (50 IU/ml) and streptomycin sulphate (50 μ g/ml). Untransfected mouse neuroblastoma cells (NB2A) and those overexpressing the full-size LPHN1 (LPH-42) were handled as described earlier⁹.

Primary human AML cells were obtained from the sample bank of the University MedicalCentre Hamburg-Eppendorf (Ethik-Kommission der Ärztekammer Hamburg, reference:

310 PV3469). Cells were incubated in IMDM medium containing 15% BIT 9500 serum 311 substitute, 100 μ M mercaptoethanol, 100 ng/ml stem cell factor (SCF), 50 ng/ml FLT3, 20 312 ng/ml G-CSF, 20 ng/ml IL-3, 1 μ M UM729 and 500 nM stemregenin 1 (SR1) as described 313 before³.

314 Primary human blood plasma samples

Blood plasma from healthy donors was generated by centrifugation of peripheral blood provided by the National Health Blood and Transfusion Service (NHSBT, UK) following ethical approval (REC reference: 16-SS-033). Primary human AML plasma samples were obtained from the sample bank of University Medical Centre Hamburg-Eppendorf (Ethik-Kommission der Ärztekammer Hamburg, reference: PV3469)⁴.

320 Western blot analysis

LPHN1 protein levels were analysed using Western blotting. β-actin staining was used to confirm equal protein loading as described previously (4, 6, 14). LI-COR goat secondary antibodies (dilution 1:2000), conjugated with fluorescent dyes, were used in accordance with manufacturer's protocol to visualise target proteins (using a LI-COR Odyssey imaging system). Western blot data were quantitatively analysed using Odyssey software and values were subsequently normalised against those of β-actin^{1,4,5}.

327 Enzyme-linked immunosorbent assays (ELISAs) and immunoprecipitation

Secreted galectin-9 was measured by ELISA using R&D Systems kits according to
manufacturer's protocol. Plasma cortisol was measured by ELISA using the Salimetrics assay
kit according to the manufacturer's protocols (Salimetrics, Suffolk, UK.)

331 For immunoprecipitation of LPHN1 fragments from human blood plasma, plates were coated332 with mouse monoclonal LPHN1 antibody following the principle and protocol described

previously⁵. This was followed by blocking with 2% BSA. Samples were then applied and 333 incubated for at least 4 h at room temperature after which plates were extensively washed 334 with Tris buffered saline with Tween-20 (TBST). Glycine-HCl pH lowering buffer (pH 2.0) 335 336 was then applied to extract the bound proteins. Extracts were mixed with equal volumes of lysis buffer (pH 7.5) and with 4× sample buffer for SDS-PAGE at a ratio of 1:3. Samples 337 were then subjected to Western blot analysis (5), using rabbit PAL1 anti-LPHN1 antibody. 338 Alternatively, PAL1 antibody was applied as detection antibody, followed by visualisation 339 using HRP-labelled anti-rabbit secondary antibody (Abcam, Cambridge, UK). 340

341 **On cell assays**

We employed a standard LI-COR on cell assay to characterise interaction of RL-1 antibody
with the surface of THP-1 cells^{4,5}.

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346 Synchrotron radiation circular dichroism (SRCD) spectroscopy

Human recombinant LPHN1 and FLRT3 were analysed, either alone or in combination with 347 each other, using SRCD spectroscopy at beamline B23, Diamond Light Source (Didcot, UK). 348 (The B23 is equipped with a highly collimated microbeam allowing the use of small aperture 349 long path length microcuvettes⁶⁻⁹). SRCD measurements were carried out using 0.01 µM 350 sample of soluble LPH-51 in a 1 cm path length cell of 3 mm aperture diameter using a 351 Module B instrument with 1 nm increment, 1 s integration time and 1.2 nm bandwidth at 23 352 °C. The cuvette capacity was 60 µl. Titration experiments were conducted as described 353 previously³ using standard far-UV measurements. 354

355 Statistical analysis

Each experiment was performed at least three times and statistical analysis was conducted using a two-tailed Student's t-test, where appropriate. Multiple comparisons were performed using an ANOVA test. Post-hoc Bonferroni correction was applied. Statistical probabilities (p) were expressed as * where p<0.05; **, p<0.01 and *** when p<0.001. Coefficient of determination (R²) was calculated using GraphPad Prism software in order to assess correlation. References 1. Sumbayev VV, Goncalves Silva I, Blackburn J, Gibbs BF, Yasinska IM, Garrett MD, et al. Expression of functional neuronal receptor latrophilin 1 in human acute myeloid leukaemia cells. Oncotarget 2016, 7: 45575-45583. 2. Volynski, KE, Meunier FA, Lelianova VG, et al. Latrophilin, neurexin, and their signaling-deficient mutants facilitate α -latrotoxin insertion into membranes but are not involved in pore formation. J Biol Chem 2000, 275: 41175-41183.

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397 Supplementary Figures

398

Cortisol facilitates immune escape of human acute myeloid leukaemia cells by inducing
latrophilin 1 expression

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411



AML THP-1 PHL

415 Supplementary figure 1. Comparative analysis of LPHN1 protein expression in primary 416 human AML, THP-1 cells and PHL. Lysates of each cell type were subjected to Western 417 blot analysis as outlined in the Materials and Methods. Images are from one experiment 418 representative of three which gave similar results. Data represent mean values \pm SEM of 419 three independent experiments; **p < 0.01 vs. AML cells.



430 Supplementary figure 2. Cell surface detection of LPHN1. Wild type NB2A cells
431 (negative control) and THP-1 cells were subjected to an on-cell assay using anti-rabbit Li-Cor
432 secondary antibody. Images are from one experiment representative of three which gave
433 similar results.

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454 Supplementary figure 3. Physiological cross-links leading to cortisol-induced 455 upregulation of LPHN1 expression in AML cells followed by facilitation of galectin-9 456 secretion in a FLRT3-dependent manner. AML is associated with a decreased blood 457 plasma glucose levels, which normally leads to upregulation of secretion of corticotropin-458 releasing hormone (CTRH) by hypothalamus. CTRH induces secretion of adrenocorticotropic 459 hormone (ACTH) by pituitary gland. Secreted ACTH upregulates cortisol production by the 460 adrenal cortex, thus leading to cortisol-induced upregulation of LPHN1 levels in AML cells.

461 Galectin-9, secreted in FLRT3-LPHN1-dependent manner attenuates anti-cancer activity of462 cytotoxic T cells (CTC) and NK cells.