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1 **Plasma low-density lipoprotein receptor-related protein 1 concentration is**
2 **not associated with human abdominal aortic aneurysm presence**

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18 deeply indebted to his contribution to this and other vascular research

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24 **What this study adds:** Polymorphisms within the LRP1 gene have been suggested to
25 contribute to AAA risk. This study demonstrates that plasma concentrations of LRP1 are
26 similar in men with and without AAA suggesting that this protein is an unsuitable marker to
27 screen at-risk populations. This study confirms that that LRP1 expression is reduced in aortic
28 biopsies collected from AAA patients compared to non-aneurysmal controls and
29 demonstrates that LRP1 inhibition reduces the ability of vascular smooth muscle cells to
30 internalise matrix metalloprotease 9 *in vitro*. These findings suggest that localised LRP1
31 dysregulation may be important in AAA pathogenesis, but is an unsuitable marker with
32 which to screen at-risk populations.

33

34 **Abstract**

35 *Objectives:* Recent genetic data suggest that a polymorphism of the low density lipoprotein
36 receptor-related protein 1 (LRP1) gene is an independent risk factor for abdominal aortic
37 aneurysm. The aims of this study were to assess whether plasma and aortic concentrations of
38 LRP1 were associated with abdominal aortic aneurysm, and to investigate the possible
39 relevance of LRP1 to abdominal aortic aneurysm pathophysiology.

40 *Design, Materials and Methods:* Three analyses were conducted. Initially, plasma LRP1
41 concentrations were measured in community dwelling men with and without abdominal
42 aortic aneurysm (n=189 and 309 respectively) using ELISA. Secondly, Western blotting
43 analyses were employed to compare the expression of LRP1 protein in aortic biopsies
44 collected from abdominal aortic aneurysm patients and non-aneurysmal post-mortem donors
45 (n=6/group). Finally, the effect of *in vitro* LRP1 blockade on MMP9 clearance by vascular
46 smooth muscle cells was assessed by zymography.

47 *Results:* Plasma LRP1 concentrations did not differ between groups of men with and without
48 abdominal aortic aneurysm (median concentration 4.56 µg/mL (IQR 3.39-5.96) and 4.43
49 µg/mL (IQR 3.44-5.84); P=0.48), and were not associated with abdominal aortic aneurysm
50 after adjusting for other risk factors (odds ratio: 1.10 (95% confidence interval: 0.91-1.32);
51 P=0.35). In contrast, LRP1 expression was ~3.4 fold lower in aortic biopsies recovered from
52 abdominal aortic aneurysm patients compared to controls (median (IQR) expression 1.72
53 (0.94-3.14) and 5.91 (4.63-6.94) relative density units; P=0.004). *In vitro* LRP1 blockade
54 significantly reduced the ability of vascular smooth muscle cells to internalise extracellular
55 MMP9.

56 *Conclusions:* These data suggest that aortic but not circulating LRP1 is down-regulated in
57 abdominal aortic aneurysm patients, and indicates a possible role for this protein in clearing
58 an aneurysm-relevant ligand.

59 **Key words:** Abdominal aortic aneurysm; Low density lipoprotein receptor related protein-1;
60 biomarker; matrix metalloprotease 9.

61

62 **Introduction**

63 Abdominal aortic aneurysm (AAA) affects ~2% men and ~1% of women aged >65 years, and
64 significantly increases the risk of mortality through aortic rupture and associated
65 cardiovascular events.^{1, 2} The aetiology of AAA remains unclear, however, inflammation,
66 extracellular matrix degeneration and vascular smooth muscle cell (VSMC) loss appear to be
67 important in the pathogenesis.^{3, 4} AAA risk factors include advanced age, male sex and
68 smoking.¹ A positive family history increases AAA risk ~2-fold, suggesting that genetic
69 factors are important in AAA development.^{5, 6} A genome-wide association study (GWAS)
70 identified a positive association of the rs1466535 major (C) allele within the low-density
71 lipoprotein receptor-related protein 1 (LRP1) gene with AAA presence (odds ratio ~1.2).⁷
72 This association appeared AAA specific and was maintained after adjusting for
73 cardiovascular risk factors leading to the suggestion that the *LRP1* rs1466535 C allele might
74 significantly contribute to AAA risk.^{8, 9} This is contradicted, however by recent data reporting
75 a positive association between AAA and the LRP1 rs1466535 T (minor) allele in a
76 geographically distinct patient population.¹⁰

77

78 LRP1 is structurally related to the low density lipoprotein receptor and comprises a 100
79 amino acid cytoplasmic domain, a membrane spanning region, and an extracellular loop with
80 4 ligand binding regions.^{11, 12} LRP1 is predominantly expressed by VSMCs and facilitates the
81 internalisation and clearance of bound ligands.¹²⁻¹⁴ Previous studies suggest that LRP1 has
82 functions beyond lipid metabolism and known LRP1 ligands include growth factors, matrix
83 metalloproteases (MMPs), protease-inhibitor complexes and extracellular matrix
84 components.^{8, 11, 12} Conditional LRP1 knockout rodent models demonstrate that LRP1
85 deficiency results in abnormal VSMC proliferation and migration, increased extracellular

86 matrix turnover and aneurysm formation within the mesenteric arteries.¹⁴⁻¹⁶ Currently, only
87 one study has directly investigated whether LRP1 expression is altered in human AAA.¹⁷ In
88 this study Chan *et al.* employed Western blotting and immunohistochemistry to demonstrate
89 significant down-regulation of the LRP1 protein in aortic biopsies of Chinese AAA patients
90 compared to non-aneurysmal controls. We hypothesised that reduced aortic LRP1
91 concentrations may result from release of this protein from the aneurysm wall, which may
92 increase circulating LRP1 concentrations in AAA patients. Here, we assess plasma LRP1 as a
93 potential AAA biomarker, confirm that aortic expression of LRP1 is significantly lower in
94 AAA biopsies compared to controls, and suggest that this protein may play a role in clearing
95 MMP9 from the aortic wall.

96

97 **Materials and methods**

98

99 Detailed materials and methods are provided in Supplementary File 1.

100

101 *Participants:* Samples collected from 3 Australian cohorts were used: 1) plasma samples
102 collected from participants of the Health In Men Study (HIMS); 2) aortic biopsies from
103 patients undergoing open surgery to repair large AAA; 3) aortic biopsies from non-
104 aneurysmal heart beating brain dead organ donors.¹⁸ The cohort characteristics and protocols
105 of the HIMS have been previously reported.¹⁸ For HIMS participants, an infra-renal aortic
106 (IRA) diameter of 30-55 mm was defined as a small AAA; and IRA diameters >55 mm were
107 considered large AAAs. Clinical information collected during the HIMS included age,
108 medical history and smoking status. For patients undergoing AAA repair, risk factors were
109 recorded as previously described (Supplementary File 1).^{19, 20} Maximum IRA diameter was
110 measured from axial computed tomography angiography (CTA) images as previously
111 described.²¹ No clinical information other than age and sex was available for organ donor
112 participants. In all instances, written informed consent and institutional ethics approval was
113 provided.

114

115 *Measurement of plasma LRP1:* Plasma LRP1 concentrations were measured using a
116 commercially available ELISA (#E91010Hu, USCN Life Sciences, China). This ELISA
117 employs antibodies against a soluble extra-cellular LRP1 immunogen suggesting suitability
118 for plasma analysis. Reported inter and intra-assay variability are <12%. Due to limited
119 plasma volumes, single measurements were taken for each patient as previously described.^{19,}

120 ²²

121

122

123 *In vitro experiments:* Human aortic VSMC (CC-2571, Lonza) were maintained in growth
124 medium in a humidified 5% CO₂ atmosphere at 37°C and passaged when 70%–80%
125 confluent. After 5 passages LRP1 blocking antibodies or isotype control antibodies (#MA1-
126 27198 and MA5-14453, respectively, Thermo Fisher Scientific, Australia) were added to the
127 culture media at concentrations of 30µg/mL. Cell culture supernatants were decanted after 24
128 hours and replaced with media containing 20ng/mL recombinant human matrix
129 metalloprotease 9 (MMP9, #911-MP-010, R and D Systems, USA). Cell culture supernatants
130 were harvested at 24 and 48 hours, and stored at -80°C. Data presented are from 3
131 independent culture experiments.

132

133 *Protein extraction and Western blotting:* Isolation of aortic proteins and western blotting was
134 performed as previously described.²³ Full-thickness human abdominal aortic samples were
135 homogenized in the presence of protease inhibitors. Samples (30 µg protein/lane) were loaded
136 onto a 10% SDS-polyacrylamide gel, electrophoresed and transferred onto a polyvinylidene
137 difluoride membrane. The membrane was cut at ~60 kDa and each half separately blocked
138 with 5% non-fat dry milk at 4°C overnight. The 60-250 kDa proteins were incubated with
139 anti-LRP1 antibody (R&D Systems MAB6360, USA), while the proteins <60 kDa were
140 incubated with anti-β actin antibody, (Abcam, UK #AB75186) for 1 hour (room
141 temperature). Membranes were washed and incubated with anti-mouse HRP- (LRP1 blot), or
142 anti-rabbit (β actin)- conjugated IgG (DakoCytomation, Denmark #P0447 and P0448
143 respectively) for 1 hour (room temperature). Membranes were washed and proteins visualised
144 with Western Lightning Chemiluminescence Reagent Plus (PerkinElmer Life Sciences,
145 USA). The relative density of the LRP1 band was standardised by dividing by mean beta-

146 actin density for the respective experimental group. Reported data refer to standardised LRP1
147 expression.

148

149 *Gelatin zymography:* Five μL of VSMC culture supernatant was loaded onto a 10%
150 polyacrylamide gel containing 1 mg/mL gelatin. Samples were electrophoresed and washed
151 twice in 2.5% (v/v) Triton X 100. Gels were incubated overnight in 50mM Tris-HCl (pH
152 8.0), 5mM CaCl_2 at 37°C and stained in 0.5% (w/v) Coomassie Blue R-250 for 30 mins. Gels
153 were destained with 40% (v/v) methanol, 10% (v/v) acetic acid, 50% (v/v) water, to visualise
154 protease activity and analysed via densitometry as above.

155

156 *Statistical analyses:* Data were analysed using SPSS v.21 (IBM) and Prism v.6 (GraphPad
157 Software Inc, San Diego, USA). Continuous variables were analysed using the Mann-
158 Whitney U test. Nominal variables were assessed using the chi-squared test. Correlations
159 were assessed using Spearman Rho analyses. The correlation between aortic LRP1
160 expression, and AAA diameter was further assessed in leave-one-out sensitivity analyses.
161 The association of plasma LRP1 with AAA presence was assessed via binary logistic
162 regression corrected for confounders identified in univariate analyses. *In vitro* data comparing
163 3 groups were assessed using one-way ANOVA applying Tukey's *post-hoc* test for multiple
164 comparisons. P values <0.05 were considered significant.

165

166 **Results**

167 *Plasma LRP1 concentrations are not associated with AAA:* LRP1 concentration was
168 measured in plasma collected from 189 men with a small AAA, and 309 non-aneurysmal

169 controls (total n=498). Participant characteristics are shown in Table 1. Participants had
170 similar age and history of dyslipidemia, diabetes and stroke. The prevalence of smoking,
171 hypertension and CHD were significantly higher in men with AAA. No differences in plasma
172 LRP1 concentration were observed between the groups Median plasma LRP1 concentration
173 was 4.56 $\mu\text{g/mL}$ (interquartile range [IQR] (3.39-5.96) for men that had AAA and 4.43
174 $\mu\text{g/mL}$ (IQR 3.44-5.84) for nonaneurysmal controls ($p=0.48$; Table 1, Fig 1A). Plasma LRP1
175 concentrations did not correlate with IRA diameter when assessing the whole population
176 (Spearman $r=0.06$, $P=0.22$; Figure 1B), or the experimental groups (for men with AAA:
177 Spearman $r=0.056$, $P=0.441$; for controls: Spearman $r=0.054$, $P=0.342$, data not shown).

178

179 Binary logistic regression was conducted to identify the major risk factors for AAA in the
180 studied population. An increase in plasma LRP1 concentration of ~ 1 std dev ($2.7 \mu\text{g/mL}$) was
181 not significantly associated with AAA presence in unadjusted analyses (odds ratio (OR): 1.07
182 (95% CI: 0.89-1.28; $P=0.476$), or in analyses adjusting for waist to hip ratio, smoking history,
183 hypertension and CHD (OR: 1.10 (95% CI: 0.91-1.32); $P=0.35$; (Supplementary Table 1). A
184 history of ever having smoked (OR: 3.44 (95% CI: 2.12-5.58), $P<0.001$), and CHD (OR: 1.94
185 (95% CI: 1.29-2.91), $P<0.01$) were independent risk factors for AAA in this population as
186 previously described (Supplementary Table 1).^{19, 22}

187

188 *Western blot analysis of aortic LRP1 expression:* As plasma LRP1 concentrations were
189 similar in AAA patients and controls, Western blotting experiments were conducted to verify
190 whether IRA LRP1 expression was lower in AAA patients compared to non-aneurysmal
191 controls (n=6/group). Patients and controls were sex-matched (67% males/group, $P=1.000$).
192 Younger AAA patients were specifically selected for this analysis to minimise age

193 imbalances between the groups although AAA patients were older than organ donors (median
194 (IQR) age 58.0 (54.8-66.6) and 45.0 (34.0-51.0) years respectively; P=0.026). Median IRA
195 diameter for the AAA patients was 65.8 (IQR: 57.3-79.2) mm. IRA measurements were not
196 available for controls. Western blotting identified a band at ~90k Da in all samples which was
197 selected for analysis as advised by the antibody manufacturer (Fig 2A and Supplementary
198 Figure 1). Standardised median aortic LRP1 expression was ~3.4-fold lower in AAA patients
199 than non-aneurysmal controls (median (inter-quartile range [IQR]) expression 1.72 (0.94-
200 3.14) and 5.91 (4.63-6.94) relative density units [RDU] respectively, P=0.004; Fig 2B). No
201 significant correlation between standardised LRP1 expression and IRA diameter in the AAA
202 patients was observed (Spearman $r=0.600$, $P=0.242$, Figure 2C). No significant correlation
203 between tissue LRP1 expression and AAA diameter was observed following leave-one-out
204 sensitivity analysis (Supplementary Table 2).

205

206 *In vitro LRP1 blockade inhibits VSMC MMP9 clearance:* The relevance of LRP1 under-
207 expression to AAA pathology remains unclear. LRP1 dysfunction has been suggested to
208 promote AAA through inefficient clearance of extracellular ligands including proteases.⁹ The
209 effect of LRP1 blockade on extracellular clearance of MMP9 by VSMCs was assessed *in*
210 *vitro*. VSMCs were exposed to LRP1 blocking antibodies, isotype control-IgG or vehicle for
211 24 hours, prior to adding recombinant MMP9. Conditioned media MMP9 activity was
212 assessed 24 and 48 hours after MMP9 addition using gelatin zymography (Supplementary
213 Figure 2).

214

215 After 24 hours, the MMP9 activity of culture supernatants was comparable between cells
216 exposed to MMP9 with anti-LRP1 antibodies, isotype control IgG, or vehicle (mean (\pm

217 standard error of the mean [SEM]) MMP9 activity: 3.69 (± 0.50), 3.85 (± 0.26) and 4.58
218 (± 0.31 RDU respectively; $P=0.270$; Figure 3A). The MMP9 activity of media from all 3
219 groups was significantly greater than negative control cells which did not receive
220 recombinant protease (all P values <0.05 ; data not shown). After 48 hours extracellular
221 MMP9 activity differed significantly between VSMCs receiving anti-LRP1 antibodies,
222 isotype control IgG, or vehicle (mean (\pm SEM): 11.27 (± 0.44), 8.29 (± 0.45) and 9.80 (± 0.75)
223 RDU respectively; $P=0.027$; Figure 3B). Between-group comparisons demonstrated that
224 conditioned media from VSMCs receiving LRP1-blocking antibodies exhibited significantly
225 higher MMP9 activity than the control IgG1-exposed cells (mean difference between groups:
226 2.97; 95% CI: 0.53-5.42; $p<0.05$). MMP9 activity from all 3 groups remained higher than the
227 negative controls at this time point (all P-values <0.05 ; data not shown).

228

229 **Discussion**

230 Recent data suggest that the LRP1 rs1466535 polymorphism may be a specific AAA risk
231 factor.^{7,9,10} A previous study has reported that LRP1 expression within the IRA is
232 significantly lower in AAA patients compared to organ donor controls.¹⁷ We and others have
233 demonstrated that proteins liberated from the aneurysm wall may enter the bloodstream and
234 act as potential biomarkers for AAA.²⁴⁻²⁶ Consequently, we hypothesised that weak aortic
235 LRP1 expression in AAA patients may increase circulating concentrations of this protein,
236 although plasma LRP1 concentrations were similar in men with (n=189) and without AAA
237 (n=309). Given this finding we attempted to verify the previous finding that LRP1 was
238 downregulated within the aorta of AAA patients. We assessed LRP1 expression in IRA
239 biopsies collected from AAA patients and organ donor controls. Currently, only one study
240 has reported significant reductions in LRP1 expression in aortic biopsies collected from AAA
241 patients compared to non-aneurysmal organ donor controls.¹⁷ Using antibodies against a
242 different LRP1 immunogen and an alternative house-keeping protein, we confirm that aortic
243 LRP1 protein expression was down-regulated in Australian patients with large AAAs
244 compared to non-aneurysmal controls. Collectively, these findings suggest that LRP1
245 dysregulation is localised to the aortic wall in AAA patients. This was assessed *in vitro* and
246 our data suggest that LRP1 blockade significantly impairs the ability of VSMCs to remove
247 the AAA relevant protease MMP9 from the surrounding media.

248

249 Bown *et al.* suggested that the LRP1 rs1466535 CC genotype is associated with increased
250 LRP1 expression (~1.2 fold) compared to the TT genotype.⁷ It should be noted that Bown
251 and colleagues analysed ascending aortic biopsies, compared to the IRA samples analysed in
252 the current study and the previous investigation by Chan *et al.*¹⁷ Gene expression patterns

253 differ between aortic regions and work is needed to assess whether the rs1466535 CC
254 genotype is associated with differential IRA LRP1 expression.²⁷ Thus, the mechanisms
255 underpinning reduced IRA LRP1 expression in AAA patients remain unclear. Chan *et al.*
256 hypothesised that LRP1 may be regulated by miR205.¹⁷ Kim and colleagues demonstrated that
257 aortic expression of miR205 is significantly increased in apolipoprotein E deficient mice
258 which developed AAA in response to angiotensin-II infusion.²⁸ Moreover administration of
259 anti-miR205 oligonucleotides to aortic endothelial cells greatly attenuated angiotensin-II
260 induced AAA development. They also showed that LRP1 expression markedly decreased in
261 response to angiotensin-II infusion, but was not rescued by *in vivo* anti-miR205
262 administration, suggesting that miR205 may exert AAA-protective effects through other
263 pathways.²⁸ Recent data also demonstrate that normocholesterolaemic mice deficient in
264 smooth muscle LRP1 (smLRP1^{-/-}) are no more susceptible to angiotensin-II induced AAA
265 than wild type controls, although the formation of large aneurysms within the ascending aorta
266 and mesenteric artery was reported.²⁹ It should be noted that normocholesterolaemic mice
267 appear more resistant to angiotensin-II induced AAA than dyslipidaemic strains and future
268 studies employing other AAA induction approaches in these mice are needed to further assess
269 the role of LRP1 in AAA formation.^{23, 30, 31}

270

271 Although there is no direct evidence of a role for LRP1 in AAA formation, studies of the
272 smLRP^{-/-} mouse have demonstrated an important role for this protein in maintaining vascular
273 integrity.^{14, 16, 32} smLRP^{-/-} mice aortas show Marfan Syndrome-like phenotypes with
274 disruption of the elastic laminae and VSMC hyperplasia, increased vessel tortuosity,
275 thickness and dilatation.^{14-16, 32} Comparable circulating cholesterol concentrations in
276 smLRP1^{-/-} and control mice suggests that the observed pathologies are independent of

277 lipoprotein metabolism.¹⁴ Intriguingly, mice deficient in both smLRP1 and LDLR appear
278 more susceptible to atherosclerosis than smLRP1⁺LDLR^{-/-} controls demonstrated by
279 widespread lesion formation following high cholesterol feeding.^{14, 16, 32} However, the
280 aetiologies of AAA and atherothrombosis appear distinct.^{20, 33} AAA is associated with VSMC
281 apoptosis and extracellular matrix degeneration, rather than cellular proliferation. MMP9, an
282 LRP1 ligand, is over-expressed in human and mouse AAA biopsies, and is implicated in
283 AAA pathogenesis.^{1, 31} The current study suggests a role for LRP1 in regulating aortic MMP9
284 concentrations evidenced by heightened proteolytic activity in supernatants from LRP1
285 compromised VSMCs. This is supported by prior data demonstrating increased MMP9
286 activity in conditioned media harvested from mouse embryonic fibroblasts treated with an
287 LRP1 antagonist, and an accumulation of MMP9 within the aortas of smLRP1^{-/-} mice.^{14, 34}
288 These findings suggest that aortic LRP1 dysregulation may exacerbate extracellular matrix
289 proteolysis, although LRP1 has potential to contribute towards multiple AAA pathogenic
290 mechanisms due to the broad spectrum of ligands for this receptor.⁹

291

292 Extracellular forms of LRP1 are thought to result from shedding of the LRP1 ectodomain.³⁵⁻³⁷
293 No association of circulating LRP1 concentration with AAA was observed suggesting that
294 low IRA LRP1 expression in AAA patients does not result from heightened extracellular
295 shedding. Immunohistochemistry data by Chan and colleagues demonstrate that LRP1 is
296 predominantly expressed within the aortic media and adventitia, and confirm that this is
297 reduced in human AAA although the cells which express LRP1 in the normal aorta were not
298 investigated in this study. LRP1 is known to be expressed by VSMCs. Thus, it is possible that
299 the loss of VSMCs typically demonstrated within the aortas of AAA patients may have

300 contributed to the down-regulation of LRP1 observed in the current study.¹⁷ Further studies
301 are required to directly assess this.

302

303 There are several limitations to the current study. Firstly, the sample sizes employed in the
304 Western blotting analyses were relatively small. As most AAAs are now managed by
305 endovascular repair aortic wall biopsies can only be obtained from the occasional patient
306 undergoing open surgery for a large AAA.¹ Similarly, ethical and practical considerations
307 prevent the collection of aortic biopsies from large numbers of control donors, however,
308 sample sizes here are similar to other published studies.^{17, 22, 38-40} Secondly the organ donor
309 controls were younger than the AAA patients and the possibility that the age imbalance may
310 have contributed to differences in aortic LRP1 expression cannot be excluded. Limited
311 clinical details were available for the organ donor controls and variation in cardiovascular
312 risk factors between groups could not be adjusted for. However, organ donors are often the
313 only viable source of healthy aortic biopsies.^{22, 41} Some previous studies have utilised
314 macroscopically healthy tissues proximal to the AAA sac as matched control tissues,
315 however these were not available for the current study.^{24, 38} A recent genome-wide expression
316 study comparing biopsies from the proximal aneurysm neck with IRA biopsies from organ
317 donors demonstrated significant reductions in LRP1 expression in the AAA patients.⁴² This
318 suggests that LRP1 dysregulation precedes macroscopic AAA dilatation. Thirdly, no plasma
319 samples were available from the AAA patients or organ donors who provided aortic biopsies.
320 Thus, IRA and plasma LRP1 concentrations could not be directly compared in these
321 individuals. Instead, plasma samples were obtained from men with screen-detected AAAs,
322 not hospital referred patients. Consequently most aneurysmal blood donors had small (<50
323 mm) AAAs. A difference in plasma LRP1 concentration may be seen when comparing
324 patients with large AAAs to healthy controls, although this was not possible to assess in this

325 study. Furthermore, the ELISA we used recognises an extracellular LRP1 immunogen
326 whereas the Western blotting antibody employed is raised against an intracellular epitope.
327 This complicates direct comparison of the two tests. Limited sample availability prevented us
328 from assessing aortic tissue extracts by ELISA to cross-validate the assay. Moreover it is
329 possible that minor differences in plasma LRP1 concentration between the groups may have
330 been masked by inter-assay variability (reported as <12%). Finally LRP1 genotype data were
331 not available for any participants in this study. The rationale for assigning participants to
332 experimental groups was therefore based on AAA.

333

334 In summary, plasma LRP1 concentrations were similar in relatively large groups of
335 community-dwelling men with and without AAA (n=189 and 309 respectively), minimising
336 the biomarker potential for this molecule. Our data support an inverse association between
337 IRA LRP1 expression and AAA presence, and suggest that LRP1 inhibition in VSMCs may
338 potentially contribute to MMP9 accumulation within the aortic wall. Studies employing
339 larger biopsy numbers and alternative methodologies such as RNA interference are required
340 to validate these findings, and to further assess the impact of reduced LRP1 expression on
341 AAA pathology.

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353

354 **Conflict of interests:** None

355

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477

478 **Table 1:** Clinical characteristics of 498 population screened men included in the plasma
479 LRP1 analyses.

Characteristic	Men without AAA (AOD <30 mm; n=309)	Men with AAA (AOD >30 mm n=189)	P-value
Infra-renal aortic diameter (mm)	19.5 (19.2-20.2)	33.7 (31.7-40.3)	<0.001
Age (years)	72 (68-75)	72 (68-75)	0.275
Waist:hip ratio	0.95 (0.91-1.00)	0.96 (0.93-1.00)	0.069
Plasma LRP1 (µg/mL)	4.43 (3.44-5.84)	4.56 (3.39-5.96)	0.476
Past history of:			
Ever smoking	196 (63.4%)	163 (86.2%)	<0.001
Hypertension	126 (40.8%)	97 (51.3%)	0.022
Dyslipidaemia	133 (43.0%)	90 (47.6%)	0.319
Diabetes	34 (11.0%)	18 (9.5%)	0.600
Stroke	19 (6.2%)	17 (9.0%)	0.234
CHD	83 (26.9%)	83 (43.9%)	<0.001

480

481 CHD: Coronary heart disease. Nominal variables are presented as numbers and valid % and
482 compared using the Chi-squared test. Continuous variables are presented as median and
483 interquartile range and compared using the Mann-Whitney U test.

484

485 **Figure legends**

486 **Figure 1:** Plasma LRP1 is not associated with AAA presence. A) ELISA analysis revealed
487 no difference in circulating LRP1 concentrations in men with AAA (n=189) and non-
488 aneurysmal controls (n=309, Mann-Whitney U test). B) No correlation was seen between
489 plasma LRP1 concentration and infrarenal aortic diameter. Non-aneurysmal controls are
490 boxed.

491

492 **Figure 2:** Western blot analysis of LRP1 expression in infra-renal aortic biopsies recovered
493 from AAA patients and non-aneurysmal organ donors. A) Immunoblot detailing relative
494 infra-renal aortic LRP1 expression in AAA patients (AAA) and controls (CTRL). B)
495 Comparisons between groups revealed standardised aortic LRP1 expression to be
496 significantly lower in AAA patients than controls. Symbols represent individual samples,
497 horizontal bars denote median and inter-quartile ranges. C) Standardised LRP1 expression
498 did not significantly correlate with infra-renal aortic diameter in the AAA patients.

499

500 **Figure 3:** The effect of *in vitro* LRP1 blockade on extracellular MMP9 activity. VSMCs
501 were cultured in the presence of recombinant MMP9 after exposure to LRP1 blocking
502 antibodies, control IgG1, or vehicle. MMP9 activity in cell culture supernatants was assessed
503 24 (A) and 48 (B) hours after MMP9 addition via gelatin zymography. Cells which did not
504 receive MMP9 are included as negative controls, but are not included in further analyses.
505 Mean \pm SEM MMP9 activity are shown for each group (n=3 for all). MMP9 activity was
506 assessed between experimental groups via one-way ANOVA, tables show mean differences

507 and 95% confidence intervals (CI) for comparisons between groups. NS: Non-significant, *:
508 $p < 0.05$ after correction via Tukey's post-hoc test for multiple comparisons.

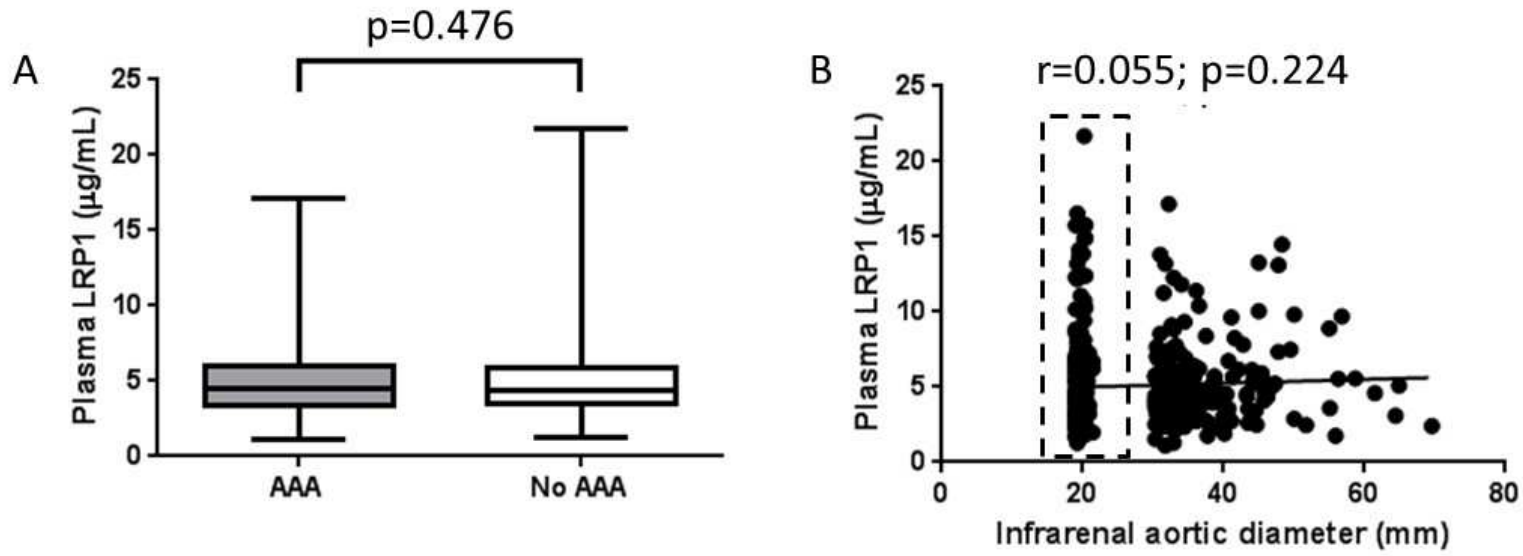


Figure 1

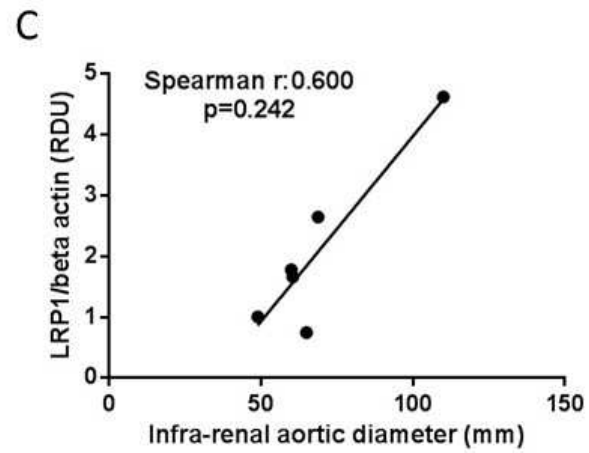
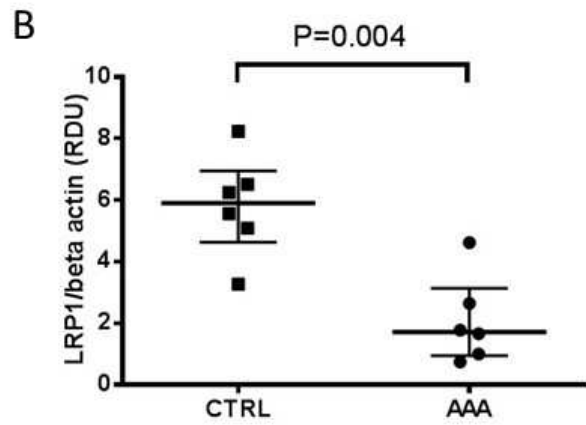
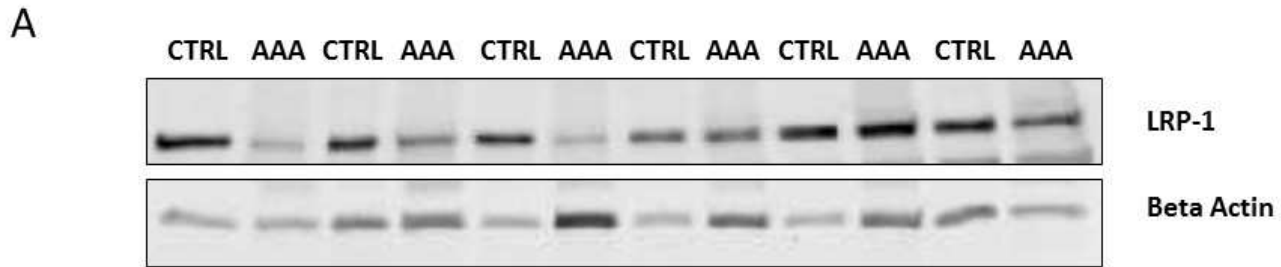


Figure 2

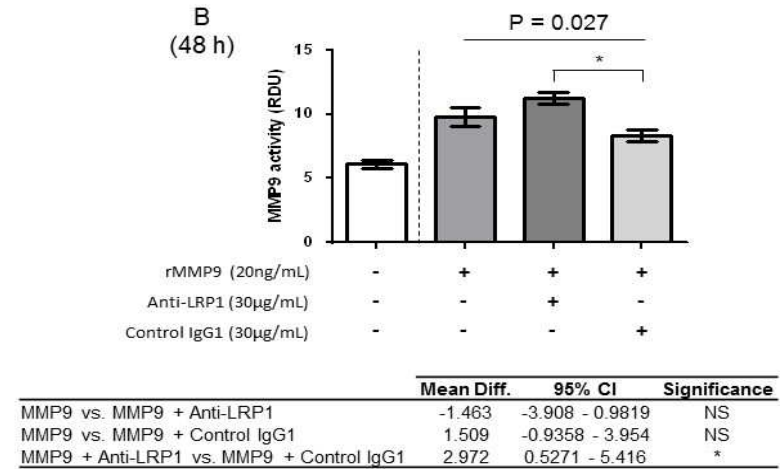
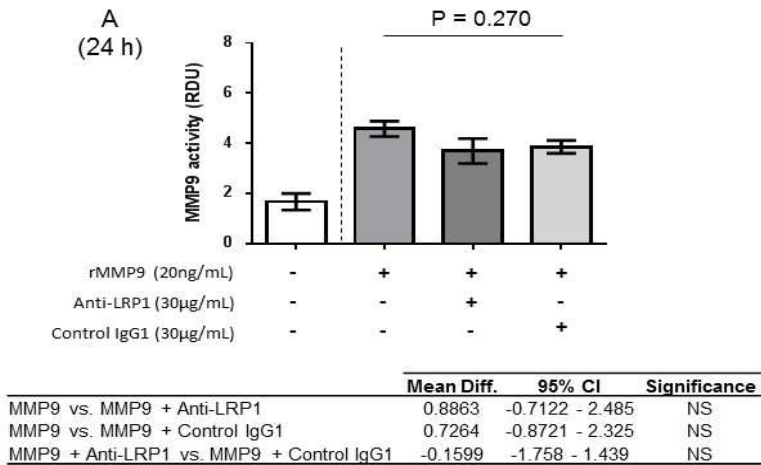


Figure 3

Moxon *et al.* Aortic wall but not plasma low-density lipoprotein receptor-related protein 1 is down-regulated in human abdominal aortic aneurysm. Supplementary material.

Supplementary File 1 – Detailed materials and methods.

Supplementary Table 1 – Independent risk factors for AAA in 498 community dwelling men screened for AAA.

Supplementary Table 2 – Leave one out sensitivity analysis assessing the correlation between aortic LRP1 expression and AAA diameter.

Supplementary Figure 1 – Raw Western blot images detailing IRA LRP1 and beta-actin expression in AAA patients and organ donor controls.

Supplementary Figure 2 – Raw zymography gels showing MMP9 activity from *in vitro* VSMC culture experiments.

SUPPLEMENTARY FILE 1 – DETAILED MATERIALS AND METHODS

Participants: Samples collected from 3 Australian cohorts were used in different stages of the study as follows: 1) aortic biopsies from patients undergoing open surgery to repair AAA, 2) aortic biopsies from non-aneurysmal heart beating brain dead organ donors, and 3) plasma samples collected from participants of the Health In Men Study (HIMS).¹⁸ For patients undergoing AAA repair risk factors were recorded as previously described.^{19, 20} Briefly, characteristics collected for each patient included sex, age, history of hypertension, diabetes, coronary heart disease (CHD) and prescribed medications. Diabetes and hypertension were defined by a prior history of diagnosis or treatment of these conditions. Smoking status was defined as having ever or never smoked. CHD diagnosis was based on a history of angina, myocardial infarction or coronary artery revascularisation. Maximum infra-renal aortic diameter was measured from axial computed tomography angiography (CTA) images using the viewer function on a Philips workstation (MxView Visualization Workstation Software, Philips Electronics) and recorded in millimeters as previously reported.²¹ No clinical information other than age and sex was available for organ donor participants. The cohort characteristics and protocols of the HIMS have been previously described.¹⁸ HIMS was a community based screening program assessing AAA which used ultrasound to measure infra-renal aortic diameter in Western Australian men aged >65 years. Maximum infra-renal aortic diameter of 30-55 mm was defined as a small AAA; and infrarenal aortic diameters >55 mm were considered large AAAs. Relevant to the current study, clinical information collected for each HIMS participant included age, medical history and smoking status. Anthropomorphic measurements including height, weight and hip and waist circumference were also recorded. Participants in the current study were selected based on blood sample availability, and completeness of clinical data. In all instances, appropriate written informed consent and

institutional ethics approval to use biological samples and clinical data for research purposes was provided.

In vitro experiments: Commercial human aortic VSMC (CC-2571, Lonza) were maintained in growth medium (DMEM (DMEM/F12, Gibco) supplemented with 10% (v/v) heat-inactivated foetal calf serum, 1% L-Glutamine and 1% penicillin/streptomycin, Gibco) in a humidified 5% CO₂ atmosphere at 37°C. Cells were passaged when 70%–80% confluent. After 5 passages VSMCs were seeded into 12 well plates at a density of 1x10⁶ cells/mL and allowed to adhere overnight. LRP1 blocking antibodies or isotype control antibodies (#MA1-27198 and MA5-14453, respectively, Thermo Fisher Scientific, Australia) were added to the culture media to final concentrations of 30µg/mL based on a previous publication using these antibodies in VSMCs.⁴³ Cell culture supernatants were decanted after 24 hours and replaced with media containing recombinant human matrix metalloprotease 9 (MMP9, #911-MP-010, R and D Systems, USA), to a final concentration of 20ng/mL. This concentration was chosen as prior data have shown this is the approximate median plasma MMP9 concentration in patients with large stable AAAs,⁴⁴ and was considered physiologically relevant. Cell culture supernatants were harvested at 24 and 48 hours, and stored at -80°C for later analysis. Data presented are from 3 independent culture experiments.

Protein extraction and Western blotting: Isolation of aortic proteins and western blotting was performed as previously described.²³ Full-thickness human abdominal aortic samples, or cultured VSMCs were homogenized in the presence of protease inhibitors to obtain protein extracts. Protein concentrations were determined using a Bradford protein assay kit (BioRad, USA). Samples (30 µg of protein per lane) were loaded onto a 10% SDS-polyacrylamide gel, electrophoresed (100 V, 90 min), and transferred (15 mA, 60 min) onto a polyvinylidene

difluoride membrane (BioRad, USA). The membrane <30 kDa was removed as mid-low molecular weight proteins were not targeted in this investigation. The remaining membrane was halved (cut at ~60 kDa) and each half separately blocked with 5% non-fat dry milk at 4°C overnight. The membrane containing the high molecular weight proteins (~60-250 kDa) was incubated with anti-LRP1 antibody (1:1000 dilution in PBSt [1 x PBS pH 7.4 + 1% (v/v) Tween 20], R&D Systems MAB6360, USA), proteins <60 kDa were incubated with anti- β actin antibody, 1:10,000 (Abcam, UK #AB75186). After 1 hour at room temperature, membranes were washed in TBSt (1 x TBS pH 7.4 + 1% (v/v) Tween 20, 3 x 10 mins) and incubated with anti-mouse HRP- (LRP1 blot), or anti-rabbit (β actin)- conjugated IgG (DakoCytomation, Denmark #P0447 and P0448 respectively) diluted 1:1,000 in PBSt for 1 hour at room temperature. Membranes were washed (2 washes in TBSt, 1 wash in TBS for 10 mins each). Protein expression was visualised with Western Lightning Chemiluminescence Reagent Plus (PerkinElmer Life Sciences, USA) and quantified using the Quantity One v4.6.7 software package (BioRad, USA). The relative density of the LRP1 band was standardised by dividing by mean beta-actin density for the respective experimental group. All presented Western blot data refer to standardised LRP1 expression.

Gelatin zymography: Five μ L of VSMC culture supernatant was loaded into each well of a 10% polyacrylamide gel containing 1 mg/mL gelatin. Samples were electrophoresed as above and then washed in 2.5% (v/v) Triton X 100 for 15 minutes. After 2 washes, gels were incubated overnight in developing buffer (50mM Tris-HCl (pH 8.0), 5mM CaCl₂) at 37°C prior to staining in 0.5% (w/v) Coomassie Blue R-250 for 30 mins. Gels were placed in destain solution (40% (v/v) methanol, 10% (v/v) acetic acid, 50% (v/v) water), to visualise areas of protease activity. Gels were analysed via densitometry as described above.

Measurement of plasma LRP1: Plasma LRP1 concentrations were measured using a commercially available ELISA according to the manufacturer's directions (#E91010Hu, USCN Life Sciences, China). This ELISA employs antibodies raised against a soluble extra-cellular LRP1 immunogen (ser4373-glu4409), suggesting suitability for plasma analysis. The manufacturer's guidelines state that the minimum detectable concentration of human LRP1 with this assay is 2.91 pg/mL and that no significant cross-reactivity with human LRP1 analogues is reported. Reported inter and intra-assay variability are both <12%. Due to limited sample availability, single measurements were taken for each patient as previously described.^{19,22}

Statistical analyses: Data were analysed using the SPSS v.21 statistical package (IBM). Plots were generated using Prism v6 software (GraphPad Software Inc, San Diego, USA). Continuous variables were compared between groups using the Mann-Whitney U test. Nominal variables were assessed using the chi-squared test. Correlation between variables was assessed using Spearman Rho analyses. The correlation between aortic LRP1 expression, and AAA diameter was further assessed in leave-one-out sensitivity analyses in which individual patients were systematically excluded. The association of plasma LRP1 with AAA presence was assessed via binary logistic regression corrected for waist:hip ratio and a history of ever smoking, hypertension or CHD based on significant, or near-significant differences between groups identified via univariate analysis. *In vitro* data comparing 3 groups were assessed using one-way ANOVA applying Tukey's *post-hoc* test for multiple comparisons. For all tests, P values <0.05 were considered significant.

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Supplementary Table 1: Independent risk factors for AAA in 498 community dwelling men screened for AAA.

Characteristic	Odds ratio	95% CI	P-value
Unadjusted			
Plasma LRP1 concentration	1.07	0.89-1.28	0.476
Adjusted for covariates			
Plasma LRP1 concentration	1.10	0.91-1.32	0.347
Waist:hip ratio	1.16	0.96-1.40	0.133
Ever smoking	3.44	2.12-5.58	<0.001
Hypertension	1.33	0.89-1.98	0.160
CHD	1.94	1.29-2.91	0.001

CHD: Coronary heart disease.

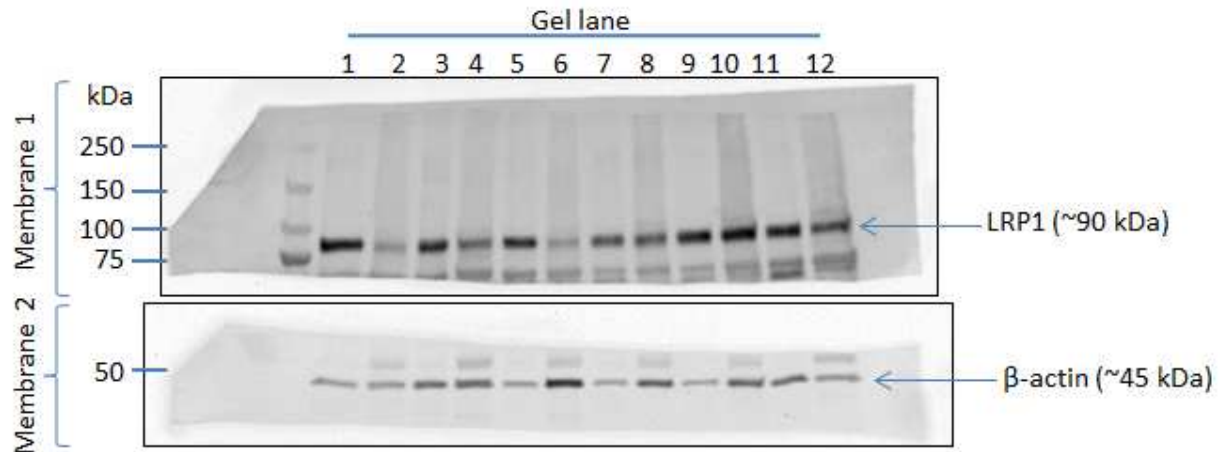
Odds ratios for plasma LRP1 concentration and waist:hip ratio refer to an increase of approximately 1 standard deviation (2.7 µg/ml and 0.06 respectively). For nominal characteristics, men with the risk factor were compared to those without.

Supplementary Table 2: Leave one out sensitivity analysis assessing the correlation between aortic LRP1 expression and AAA diameter.

Patient excluded*	AAA diameter (mm)	LRP1 expression (RDU)[†]	Spearman's r	P-value
1	49.0	1.01	0.600	0.350
2	60.0	1.78	0.700	0.233
3	65.0	0.75	0.900	0.083
4	60.5	1.67	0.700	0.233
5	110.0	4.62	0.300	0.683
6	68.9	2.65	0.300	0.683

* Patient ID is arbitrary

[†] Refers to LRP1 expression measured by Western blot (normalised to mean beta-actin expression for the whole group); RDU: Relative Density Units.



Supplementary Figure 1: Raw images of western blots detailing LRP1 (membrane 1) and β -actin (membrane 2) expression in aortic biopsies recovered from human organ donors (lanes 1, 3, 5, 7, 9 and 11) and AAA patients (lanes 2, 4, 6, 8, 10 and 12). Each lane contains 30 μ g protein and lane numbering is uniform for both membranes. After western blotting, membranes were cut between the 75 and 50 kDa markers, and probed with antibodies against LRP1 (membrane 1) or β -actin (membrane 2). As this experiment did not target medium-small (<35 kDa) proteins, portions of the membranes spanning this region were discarded.

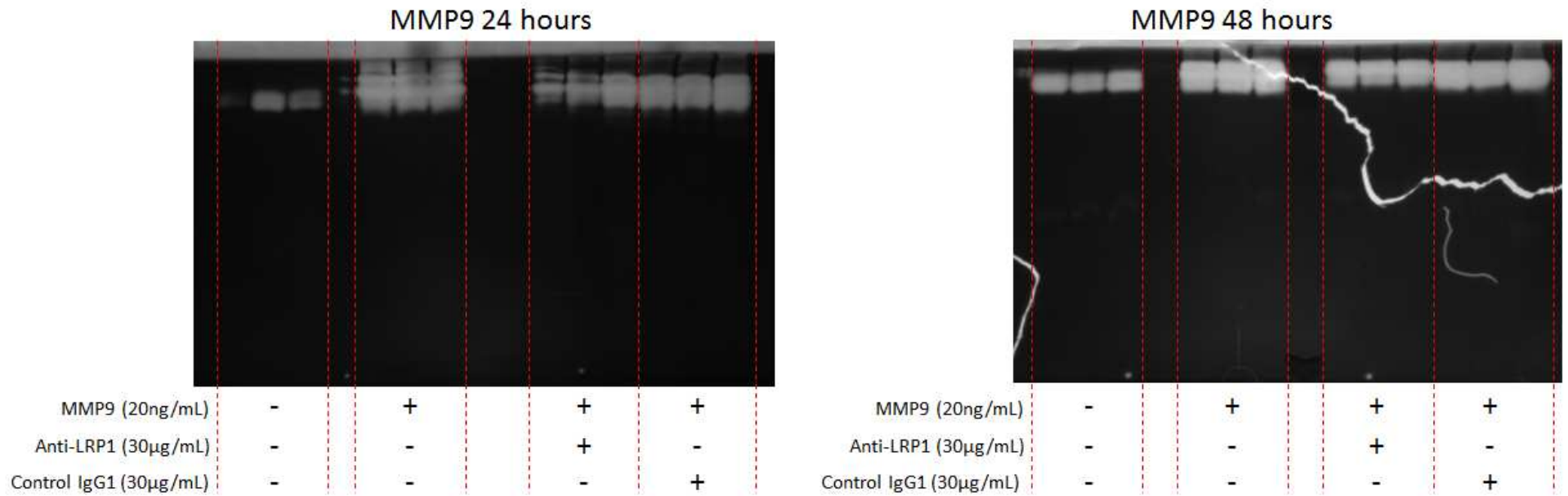
After incubating in primary and secondary antibodies, immune complexes were visualised via chemiluminescence. Membranes were imaged separately (exposure times: 250 seconds (membrane 1) and 300 seconds (membrane 2)), and relative expression of each protein compared between samples via densitometry. No inter-membrane comparisons were made.

The Western blotting antibody (R&D Systems MAB6360) was selected after considering the following key factors:

- 1) The manufacturer specifically advises that the antibody is recommended for use in Western blotting analysis. Moreover, quality control data specifically state that no-cross reactivity was observed with LRP1 homologues. Collectively this suggested that the antibody displays high specificity for LRP1 under the experimental conditions employed for the current study.
- 2) The antibody is raised against an immunogen within the LRP1 intra-cellular domain, providing scope to specifically assess endogenous LRP1 expression within the aortic wall.
- 3) The antibody has been used in a recent high-impact publication,¹ indicating that it has been validated within the wider scientific community.

Reference:

- 1) Hayashi H, *et al.* (2012). A potential neuroprotective role of apolipoprotein E-containing lipoproteins through low density receptor-related protein 1 in normal tension glaucoma. *J Biol Chem* 287(3) 25395-406



Supplementary Figure 2: Zymography gels showing MMP9 activity in conditioned media collected from VSMCs 24 and 48 hours after adding MMP9 to the culture. Details of reagents added to each culture are shown.