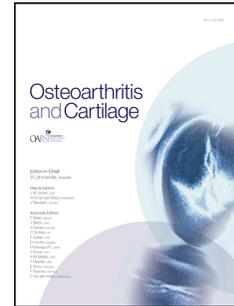


# Accepted Manuscript

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Sally Roberts, Helena Evans, Karina Wright, Louw van Niekerk, Bruce Caterson,  
James B. Richardson, Karadi Hari Sunil Kumar, Jan Herman Kuiper



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**ADAMTS-4 ACTIVITY IN SYNOVIAL FLUID AS A BIOMARKER OF  
INFLAMMATION AND EFFUSION**

<sup>1,2</sup>Sally Roberts, <sup>1</sup>Helena Evans, <sup>1,2</sup>Karina Wright, <sup>3</sup>Louw van Niekerk, <sup>4</sup>Bruce Caterson,  
<sup>1,2</sup>James B Richardson, <sup>1</sup>Karadi Hari Sunil Kumar and <sup>1,2</sup>Jan Herman Kuiper,  
<sup>1</sup>Robert Jones & Agnes Hunt Orthopaedic Hospital NHS Foundation Trust, Oswestry,  
Shropshire & <sup>2</sup>Keele University, Keele, Staffordshire, UK; <sup>3</sup>Northallerton Hospital,  
Yorkshire and <sup>4</sup>School of Bioscience, Cardiff University, UK.

**email addresses:**

H Evans: helena.evans@rjah.nhs.uk

K Wright: karina.wright@rjah.nhs.uk

Lv Niekerk: shelly.cameron@stees.nhs.uk

B Caterson: caterson@cardiff.ac.uk

JB Richardson: james.richardson@nhs.net

S Kumar: drkhskumar@yahoo.com

JH Kuiper: jan.kuiper@nhs.net

**Corresponding Author:**

Sally Roberts

ARC/TORCH Building & ISTM (Keele University)

Robert Jones & Agnes Hunt Orthopaedic Hospital

Oswestry, Shropshire, SY10 7AG, UK

tel -44 1691 404664

email: sally.roberts@rjah.nhs.uk

**ABSTRACT****Objective:**

To evaluate the potential of ADAMTS-4 (aggrecanase -1) activity in synovial fluid as a biomarker of knee injury and joint disease.

**Design:**

We have measured ADAMTS-4 activity in the synovial fluid of 170 orthopaedic patients with different degrees of joint pathology, using a commercial ADAMTS-4 fluorescence resonance energy transfer (FRET) substrate assay. Patients were classified at arthroscopy as (i) macroscopically normal, (ii) with an injury of the meniscus, anterior cruciate ligament or chondral/osteochondral defects or (iii) with osteoarthritis, and the influence of independent factors (age, patient group, effusion and synovial inflammation) on ADAMTS-4 activity levels was assessed.

**Results:**

In most patients (106/170) ADAMTS-4 activity was undetectable; ADAMTS-4 ranged from 0-2.8ng/mL in synovial fluid from patients with an injury, 0-4.1ng/mL in osteoarthritic patients and 4.0-12.3ng/mL in patients with large effusions. Four independent variables each significantly influenced ADAMTS-4 activity in synovial fluid (all  $p < 0.001$ ): age (concordance=0.69), presence of osteoarthritis (OA) (concordance=0.66), level of effusion (concordance=0.78) and inflammation (concordance=0.68). Not only did effusion influence the amount of ADAMTS-4 activity most strongly, but it also did this in an ordered manner ( $p < 0.001$ ).

**Conclusions.**

The main finding of this study is that ADAMTS-4 levels in synovial fluid are most strongly correlated with inflammation and severity of effusion in the knee. Further study is required to determine if it could provide a useful tool to aid clinical diagnoses, indicate treatment, to

monitor progression of joint degeneration or osteoarthritis or alternatively the success of treatment.

**KEYWORDS:** ADAMTS-4 activity, synovial fluid, inflammation, effusion, knee osteoarthritis

**RUNNING TITLE:** ADAMTS-4 activity as a biomarker

## 1 INTRODUCTION

2 The lack of valid biomarkers with a good relationship to joint structural pathology and symptomatic  
3 disease in individual patients is suggested as being partly responsible for the slow pace of  
4 innovation in developing novel, effective treatments for osteoarthritis<sup>1</sup>. Osteoarthritis (OA) of the  
5 knee can develop secondary to several disorders or injury in the joint and manifests as alteration of  
6 the joint structure, with progressive degradation of any or all of the tissues within it, such as  
7 cartilage, menisci and ligaments, as well as inflammation of the synovium and changes to the  
8 subchondral bone. Measuring biomarkers in fluids proximal to the site of the pathology, such as the  
9 synovial fluid in the knee, can be both more informative about the disease state in that particular  
10 joint and more sensitive, by virtue of their higher concentration than, for example, in blood or  
11 urine<sup>2,3</sup>.

12  
13 The best candidates for biomarkers in OA are suggested to be structural molecules or enzymes  
14 linked to cartilage, bone or synovium degradation<sup>4</sup>. One of the earliest and most striking  
15 biochemical changes to articular cartilage following injury or in degenerative joint diseases, such as  
16 OA, is degradation and loss of aggrecan. Matrix metalloproteinases (MMPs) and aggrecanases (or  
17 A Disintegrin And Metalloprotease with Thrombospondin motifs (ADAMTS)) are the enzymes  
18 attributed with degrading the majority of the aggrecan. The most common 'pathological'  
19 ADAMTS-generated cleavage site in the aggrecan core protein is between the <sup>392</sup>glutamate and the  
20 <sup>393</sup>alanine bond in the interglobular domain (reviewed in <sup>5</sup>). Many studies have been undertaken to  
21 develop and trial measuring the presence of these 'neo-epitopes' generated by enzyme activity to  
22 assess their potential as biomarkers with an ARGS-aggrecan assay showing considerable promise  
23 when measured in blood or synovial fluid<sup>6</sup>.

24

1 ADAMTS-5 (aggrecanase-2) is the predominant member of the ADAMTS family in mice models  
2 of degenerative joint disease or OA. The situation in humans is less clear and ADAMTS-4  
3 (aggrecanase-1) is likely to be a significant player<sup>5</sup>. In this study we have used a commercially  
4 available assay that can detect ADAMTS-4 activity independently of any contribution from  
5 ADAMTS-5. We have measured the extent of ADAMTS-4 activity in synovial fluid from a cohort  
6 of heterogeneous patients, typical of those attending an orthopaedic outpatient clinic with different  
7 degrees of joint pathology, ranging from macroscopically normal to end stage OA. Our objective  
8 was to investigate if measuring ADAMTS-4 activity could provide a useful tool to aid the clinician.

## 10 **METHOD**

### 11 **(i) Patient samples**

12 Synovial fluid was collected from patients presenting with clinical symptoms who were undergoing  
13 a routine diagnostic arthroscopy of the knee and consented to take part in this study (99\61\RJ  
14 approved by Shropshire Research Ethics Committee). Patients were classified into 3 groups  
15 according to their appearance at arthroscopy as assessed by the treating surgeon:

- 16 (i) those with macroscopically normal knees and no obvious abnormality,
- 17 (ii) those with injury of the meniscus, anterior cruciate ligament or chondral/osteochondral lesions  
18 and
- 19 (iii) those with OA (i.e. with features, such as the presence of osteophytes and fragmenting articular  
20 cartilage in one or more compartments of the knee (Outerbridge Grade 3<sup>7</sup>) or exposed subchondral  
21 bone (Outerbridge Grade 4).

22 The presence of synovial inflammation was noted when the synovium was more reddened, swollen  
23 and convoluted than one would expect in a normal knee joint and the extent of joint effusion (none,  
24 small, moderate or large) was also assessed, using the sweep and patella tap tests<sup>8</sup>. A small effusion  
25 was recorded if the sweep test was positive but the patella tap test negative, a moderate effusion was

1 recorded if the sweep and patella tap tests were positive and a large effusion was recorded if sweep  
2 and patella tap tests were negative due to over-distention of the joint with effusion.

3  
4 The synovial fluid was collected by injecting 20mLs of 0.9% saline into the synovial cavity prior to  
5 arthroscopy; the knee was flexed and extended 20 times to allow mixing of the saline with the joint  
6 fluid. A needle was then reinserted into the knee and the synovial fluid aspirated; in the laboratory it  
7 was centrifuged for 15 minutes at 3000g and the supernatant stored at -80°C until use.

8  
9 Plasma was collected at the same time and a dilution factor of the synovial fluid (SF) was obtained  
10 by measuring the ratio of urea in SF:plasma (normally ~0.9 SF:serum)<sup>9</sup>. Urea was measured with a  
11 colorimetric assay (QuantiChrom™ Urea assay Kit (BioAssay Systems DIUR-500)).

12  
13 (ii) **ADAMTS-4 activity** was measured in synovial fluid with the Sensolyte™ 520 ADAMTS-4  
14 assay kit (Anaspec Inc) which uses a substrate for fluorescence resonance energy transfer (FRET),  
15 5-FAM and TAMRA as the donor-acceptor pair. This acts as a substrate for active ADAMTS-4,  
16 which cleaves it into two separate fragments resulting in an increase of 5-FAM fluorescence which  
17 can be monitored at excitation and emission of 490nm and 520nm, respectively. The ADAMTS-4  
18 assay was carried out according to the manufacturer's instructions. Fifty microlitres of each sample  
19 of SF lavage were pipetted into wells of a 96-well black microplate (Costar®, Corning Life  
20 Sciences). In addition, a range of concentrations (2-10ng) of human recombinant ADAMTS-4 were  
21 pipetted into other wells to provide a standard curve to relate ADAMTS-4 activity in the SF  
22 samples to ng/mL of enzyme. There was also a well containing assay buffer only, to provide a  
23 substrate control. Fifty microlitres of 5-FAM fluorescence reference standards (70-2500nM) were  
24 set up to calibrate the plate reader and act as an indicator of the amount of final product of the  
25 enzymatic reaction. The substrate solution (5-FAM/5-TAMRA) was added after pre-incubation at

1 37°C and the fluorescence intensity was measured following 60 minutes incubation at 37°C.  
2 Relating the amount of fluorescence intensity from each sample to that of the standard curve  
3 obtained from the recombinant ADAMTS-4 and multiplying it by the appropriate dilution factor for  
4 each SF sample, calculated the amount of ADAMTS-4 protein in each sample.

5

### 6 (iii) **Statistics**

7 The lower limit of detection (LLOD) and lower limit of quantification (LLOQ) of ADAMTS-4  
8 were calculated from the standard deviation of blank samples and the slope of the calibration curve.  
9 QQ-plots were used to assess the distribution of continuous variables; age was normally distributed  
10 but ADAMTS-4 level was not. ADAMTS-4 levels were therefore summarised using medians and  
11 quartiles and compared between groups using non-parametric tests whereas parametric tests were  
12 used for age. Fisher's exact test was used to compare distributions in categorical variables (e.g.  
13 presence of effusion by patient classification). ADAMTS-4 levels below the LLOD were classified  
14 as having 0ng/mL of ADAMTS-4. We used a censored or Tobit regression model<sup>10</sup> to investigate  
15 the influence of the various independent factors (age, patient group, effusion and inflammation) on  
16 ADAMTS-4 levels, with the left limit set at the LLOD. This method requires a normal distribution  
17 for ADAMTS-4 levels, which was achieved by a logarithmic transformation, and minimises  
18 potential bias from the presence of a LLOD compared to traditionally used methods such as  
19 imputing half the LLOD for values below the LLOD<sup>10,11</sup>. The association between the independent  
20 factors (e.g. disease/injury, effusion and the presence of an inflamed synovium) and ADAMTS-4  
21 levels was assessed by the concordance  $c$  (a generalisation of the area under the curve (AUC) of a  
22 receiver operating characteristic (ROC) curve) and Nagelkerke's  $R^2$ , with 95% confidence intervals  
23 calculated on the basis of 999 bootstrap samples.

24

1 All statistical analysis was performed using R vs 3.0.3 (The R Foundation for Statistical  
2 Computing) using the packages “censReg” and “rms”. A p-value of 0.05 or below was assumed to  
3 denote statistical significance. In cases of multiple pairwise tests, p-values were adjusted using  
4 Holm’s procedure to maintain the 0.05 alpha-level.

5

## 6 **RESULTS**

7 Samples were collected from 170 patients, presenting with symptoms for varying times, ranging  
8 from 1 month to 16 years. The mean age of the macroscopically normal group was 28.6 years  
9 (8.5SD; n=22), the injury group 33.2 years (10.3SD; n=96) and the OA group 48.7 years (16.9SD;  
10 n=10). A number of patients combined an injury and OA (mean age 48.9 years  $\pm$ 11.6SD; n=31).  
11 Mean age differed significantly between patients with OA (alone or combined with injury) and  
12 those with ‘macroscopically normal’ joints (both  $p<0.001$ ) or injured joints (both  $p<0.001$ , all  
13 pairwise t-tests). Eleven patients were not classified. Effusions were present in 6% of ‘normal’  
14 joints, 40% of those with injuries, 71% of those with OA and 64% of those with injuries and OA.  
15 Effusions were less common in ‘macroscopically normal’ joints than any of the other types  
16 ( $p<0.05$ ) but otherwise the incidence rates did not differ between the groups ( $p>0.25$ , all pairwise  
17 Fisher’s exact tests). Inflammation was noted in the knee joints of 25% of patients categorised as  
18 ‘macroscopically normal’, 54% of the injury group, 67% of the OA group and 83% of those with  
19 injuries and OA. These rates were higher in joints with OA and injury compared to ‘normal’ joints  
20 or those with only injury ( $p<0.05$ ), otherwise no differences were found ( $p>0.09$ , all pairwise  
21 Fisher’s exact tests).

22

23 The lower limit of detection (LLOD) and lower limit of quantification (LLOQ) of the ADAMTS-4  
24 assay were 0.14ng/mL and 0.41ng/mL, respectively. In most patients ADAMTS-4 activity did not  
25 reach detectable levels (106 out of 170). It was detectable in only 5% of samples from the ‘normal’

1 group, in 33% of the injury group, in 60% of the OA group and in 67% of the combined injury and  
2 OA group. These rates were lower in the 'normal' group compared to each of the three others  
3 ( $p < 0.03$ ) and lower in the injury group compared to the combined injury and OA group ( $p = 0.004$ ),  
4 but otherwise no differences were found ( $p > 0.32$ , all pairwise Fisher's exact tests). The measured  
5 level of ADAMTS-4 ranged between 0.2 and 34.9 ng/mL SF (with a mean dilution factor for the SF  
6 samples of  $3.6 (\pm 2.2, \text{range } 1.1-12.1)$ ). The median level of ADAMTS-4 was 0 ng/mL in the  
7 'normal' group (with one patient having a detectable amount of 0.3 ng/mL), 0 ng/mL in the injury  
8 group (range 0 to 29.8, with 32 patients having detectable levels), 1.2 ng/mL in the OA samples  
9 (range 0 to 11.5, with 6 patients having detectable levels) and 1.3 ng/mL in the combined injury and  
10 OA group (range 0 to 15.4, with 21 patients having detectable levels). There was significantly less  
11 ADAMTS-4 in the 'normals' than any of the other 3 patient groups and the injury group had less  
12 than the combined injury and OA group (both  $p < 0.005$ ; pairwise Mann-Whitney tests).

13  
14 Further analysis was simplified by separating the patient group qualifier into two: having OA  
15 (No/Yes) and having an injury (No/Chondral/Other; Table 1). No significant interaction effect of  
16 injury and OA on ADAMTS-4 levels was found ( $p = 0.13$ ; Tobit regression), justifying their further  
17 analysis as separately additive.

18  
19 Four independent variables each significantly influenced ADAMTS-4 levels (all  $p < 0.001$ ; Table 1,  
20 univariable analysis), namely age (concordance=0.69; Nagelkerke's  $R^2 = 0.10$ ), having OA  
21 (concordance=0.66; Nagelkerke's  $R^2 = 0.11$ ), effusion level (concordance=0.78; Nagelkerke's  
22  $R^2 = 0.31$ ) and presence of an inflamed synovium (concordance=0.68; Nagelkerke's  $R^2 = 0.11$ ).  
23 Effusion had the best predictive value, with ADAMTS-4 levels rising in an ordered fashion with  
24 increased effusion ( $p < 0.001$ , Jonckheere test; Fig. 1). In all, 14% of patients with no effusion, 36%  
25 of patients with a small effusion, 71% with a moderate effusion and 89% of those with a large

1 effusion had detectable levels of ADAMTS-4. The significance levels from the univariable Tobit  
2 regression and conventional non-parametric tests were in complete agreement (Table 1).

3  
4 The Tobit regression allows simultaneous investigation of multiple factors. Using all variables (full  
5 model; Table 1) suggested that three independent variables simultaneously significantly influenced  
6 ADAMTS-4 levels, namely age, level of effusion and having an inflamed synovium (reduced  
7 model, Table 1).

## 8 9 **DISCUSSION**

10 Several international efforts have been and are addressing the challenge of developing reliable,  
11 sensitive and specific biomarkers for OA. Examples are those by the European Society for Clinical  
12 Aspects of Osteoporosis and Osteoarthritis (ESCEO)<sup>4</sup> and the Osteoarthritis Research Society  
13 International – Federal Drugs Agency (OARSI-FDA) initiative ([http://oarsi.org/education/oarsi-  
14 resources/fnih-osteoarthritis-biomarkers-consortium-project](http://oarsi.org/education/oarsi-resources/fnih-osteoarthritis-biomarkers-consortium-project)). Many groups have investigated the  
15 aggrecanase-generated neoepitope ARGS and assays are now available with great sensitivity<sup>12</sup>  
16 (down to 0.025pmol/mL<sup>6</sup>). This neoepitope can be measured not only in synovial fluid, but also in  
17 serum, plasma and urine, although results are less reliable in urine<sup>13;6</sup>. These assays, however, will  
18 not discriminate between the different forms of aggrecanase or ADAMTSs. The assay used in the  
19 present study measured activity of ADAMTS-4 (aggrecanase 1), with negligible cleavage of the  
20 FRET substrate by other members of the aggrecanase family, such as ADAMTS-1 and ADAMTS-  
21 5. Another ADAMTS-4 assay, using a fluorescent ‘turn-on’ peptide conjugated to gold  
22 nanoparticles, demonstrated differential levels in acute and chronic joint injuries<sup>14</sup>.

23  
24 Our data suggests that effusion level, presence of synovial inflammation and age are the strongest  
25 determinants of level of ADAMTS-4, with injury or OA being less important. Mechanical loading  
26 is known to influence protease production, with the type of load switching production between

1 MMPs and aggrecanases, e.g. in rat intervertebral disc cells<sup>15</sup>. In a murine model of OA it has been  
2 shown that gene expression of proteases, including ADAMTS-4 and -5, is both rapid and highly  
3 mechanosensitive<sup>16</sup>. This could explain the particularly strong relationship which we saw between  
4 effusion (and so swelling and increased pressure) in the joint and levels of ADAMTS-4 activity.  
5 Since ADAMTS-5 activity has not been assessed in the present study, it would be very interesting  
6 to measure this in the same or similar patient groups to determine if it also is increased with  
7 effusion.

8  
9 The relative levels of MMP and aggrecanase production in a joint may be important not only to how  
10 the joint tissues degrade but also to the individual's ability to heal that injury and to their likelihood  
11 of developing OA. The DBA/1 strain of mice, which shows superior healing of injuries compared to  
12 the C57BL/6 strain, have less aggrecanase- but more MMP-induced aggrecan degradation than the  
13 C57BL/6 mice. In addition, the DBA/1 mice did not go on to develop subsequent signs of OA seen  
14 in the C57BL/6 mice<sup>17</sup>.

15  
16 If ADAMTS-4 proves useful as a biomarker, further information on the aetiopathogenesis of  
17 degenerative joint disease(s) may be possible by measuring a differentially spliced variant of  
18 ADAMTS-4 which appears to be produced predominantly by synovial cells, particularly in OA<sup>18</sup>.  
19 Certainly differentiating the tissue source is important for biomarkers as ADAMTS-4 has been  
20 shown to be involved in atherosclerosis and is produced by monocytes from patients with acute  
21 coronary syndrome<sup>19</sup>. In addition, identification that ADAMTS-4 is a key player can help identify  
22 patients who might benefit from specific ADAMTS-4 inhibitors which are being developed. In  
23 terms of the BIPED classification of biomarkers<sup>20</sup> (Burden of disease, Investigative, Prognostic,  
24 Efficacy of intervention or Diagnostic), evidence from this study suggests that ADAMTS-4 may  
25 prove useful for investigating the aetiopathogenesis of disease, making a diagnosis of joint effusion

1 or inflammation and perhaps assessing the burden of the disease. Conversely, our results also  
2 suggest that the extent of joint effusion might serve as a proxy for ADAMTS-4 levels in the joint  
3 fluid. The low concordance and large influence of confounding factors suggest ADAMTS-4 is  
4 unlikely to be suitable for OA diagnosis.

5  
6 There are, of course, several limitations to this study. One such is the way we related the relative  
7 fluorescent units created by the ADAMTS-4 in the SF samples to protein levels (ng/mL). We made  
8 the assumption that the kinetic curve would be the same for the recombinant ADAMTS-4 as it  
9 would be for the ADAMTS-4 in the biological samples, which of course it may not be. Different  
10 truncations and portions of the enzyme give different kinetics<sup>18</sup>. Variation in ADAMTS-4 structure,  
11 such as has been seen in OA joints, warrants further investigation in this matter in future studies.  
12 Another limitation is that we have only assessed ADAMTS-4 activity levels in comparison to a  
13 limited number of variables and observations and there are always more which could be made. For  
14 example, it would be interesting to determine if there was a relationship with the duration of  
15 patients' symptoms. A further problem with the study is missing data. Ideally, we would have full  
16 data on all patients but as in most 'real world' clinical studies this was not possible and for some  
17 patients information is partly lacking (e.g. 11 patients (6.5%) were not classified by the surgeon into  
18 1 of the 3 categories at the time of the arthroscopic assessments). We believe that this data is  
19 missing completely at random as a consequence of busy clinics and therefore 'missingness' should  
20 not add bias to the assessments.

21  
22 In conclusion, identification and validation of reliable, sensitive biomarkers for injured or  
23 degenerate joints remain important goals, not only for patient diagnosis and prognosis, but also for  
24 furthering our understanding of disease progression and indicating therapies which can be applied at

1 a stage when the disease can be alleviated or modulated. We believe that measuring the level of  
2 ADAMTS-4 shows promise as a potentially useful biomarker and is worthy of further study.

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6 Council (G0800248) for their support in carrying out this project.

#### 8 **Author contributions**

9 **Sally Roberts:** Conception and design, analysis and interpretation of the data, drafting of the article,  
10 critical revision of the article for important intellectual content, final approval of the article,  
11 obtaining of funding, administrative, technical or logistic support, collection and assembly of data.

12 She takes responsibility for the integrity of the work as a whole (sally.roberts@rjah.nhs.uk)

13 **Helena Evans:** Analysis and interpretation of the data, drafting of the article, final approval of the  
14 article, technical or logistic support, collection and assembly of data.

15 **Karina Wright:** Analysis and interpretation of the data, drafting of the article, final approval of the  
16 article.

17 **Louw van Niekerk:** Drafting of the article, final approval of the article, provision of study material  
18 or patients, collection and assembly of data.

19 **Bruce Caterson:** Conception and design, drafting of the article, critical revision of the article for  
20 important intellectual content, final approval of the article.

21 **James B Richardson:** Drafting of the article, final approval of the article, provision of study  
22 material or patients, collection and assembly of data.

23 **Karadi Hari Sunil Kumar:** Drafting of the article, final approval of the article, provision of study  
24 material or patients, collection and assembly of data.

25 **Jan Herman Kuiper:** Analysis and interpretation of the data, drafting of the article, final approval  
26 of the article, statistical expertise.

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Table 1. Univariable and Multivariable predictors of ADAMTS-4 in synovial fluid of the knee

	n <sup>1</sup>	Median ADAMTS-4 ng/mL (IQR)	p-value <sup>2</sup>	Coefficient <sup>3</sup> (95% CI)	Concordance c (95% CI)	Nagelkerke's R <sup>2</sup> (95% CI)	p-value <sup>4</sup>
<b>Univariable analysis</b>							
Age	170			0.08 (0.04-0.13)	0.69 (0.62-0.78)	0.10 (0.03-0.19)	<0.001
OA	159		<0.001				
No	118	0.0 (0-0.3)		2.4 (1.2-3.6)	0.66 (0.59-0.73)	0.11 (0.04-0.22)	<0.001
Yes	41	1.3 (0-4.1)					
Injury	160		0.10				
None	32	0 (0-0)			0.58 (0.52-0.65)	0.03 (0.00-0.12)	0.10
Chondral	95	0 (0-2.2)		1.7 (0.2-3.3)			
Other	33	0 (0-2.8)		1.4 (-0.4-3.3)			
Effusion	122		<0.001 <sup>5</sup>				
No	71	0.0 (0.0-0.0)			0.78 (0.69-0.86)	0.31 (0.17-0.49)	<0.001
Small	28	0.0 (0.0-0.75)		1.8 (0.4-3.2)			
Moderate	14	1.7 (0.2-3.4)		3.6 (1.9-5.3)			
Large	9	8.5 (4.0-12.3)		5.4 (3.4-7.3)			
Inflammation	144		<0.001				
No	62	0.0 (0.0-0.0)			0.68 (0.60-0.76)	0.11 (0.03-0.24)	<0.001
Yes	82	0.2 (0.0-2.8)		2.4 (1.1-3.6)			
<b>Multivariable analysis</b>							
Full model	104				0.88 (0.83-0.96)	0.46 (0.36-0.66)	
Age					0.72 <sup>6</sup>	0.13 <sup>6</sup>	0.07
OA					0.65	0.10	0.64
Injury					0.54	0.01	0.86
Effusion					0.79	0.34	<0.001
Inflammation					0.70	0.15	0.04
Reduced model	106				0.88 (0.83-0.95)	0.46 (0.34-0.63)	
Age				0.05 (0.01-0.09)	0.72 <sup>6</sup>	0.13 <sup>6</sup>	0.02
Effusion					0.79	0.34	<0.001
No							
Small				1.7 (0.4-3.0)			
Moderate				3.3 (1.7-4.8)			
Large				4.4 (2.6-6.2)			
Inflammation					0.71	0.16	0.006
No							
Yes				1.8 (0.5-3.0)			

Notes 1) Number of patients for each analysis, which vary due to data missingness 2) p-values from univariate non-parametric statistical analyses. 3) Coefficients from censored regression. 4) p-values from censored regression. 5) p-value from Jonckheere trend test. 6) Concordance and Nagelkerke's R<sup>2</sup> values for individual predictors in the full and reduced models are the univariable values when calculated using each model's dataset. They are therefore NOT partial concordance or R<sup>2</sup> coefficients. (IQR: interquartile range; CI: confidence intervals)

Figure 1.

Levels of ADAMTS-4 (ng/mL) activity as a function of effusion size, shown as a boxplot.

The thick horizontal lines represent the medians, the boxes represent upper and lower quartiles and the fences indicate the range up to 1.5 times the interquartile range below or above the lower or upper quartiles. Values higher than 1.5 times the interquartile range above the upper quartile are shown as open circles. The lower limit of detection of 0.14 ng/ml is shown as a horizontal line.

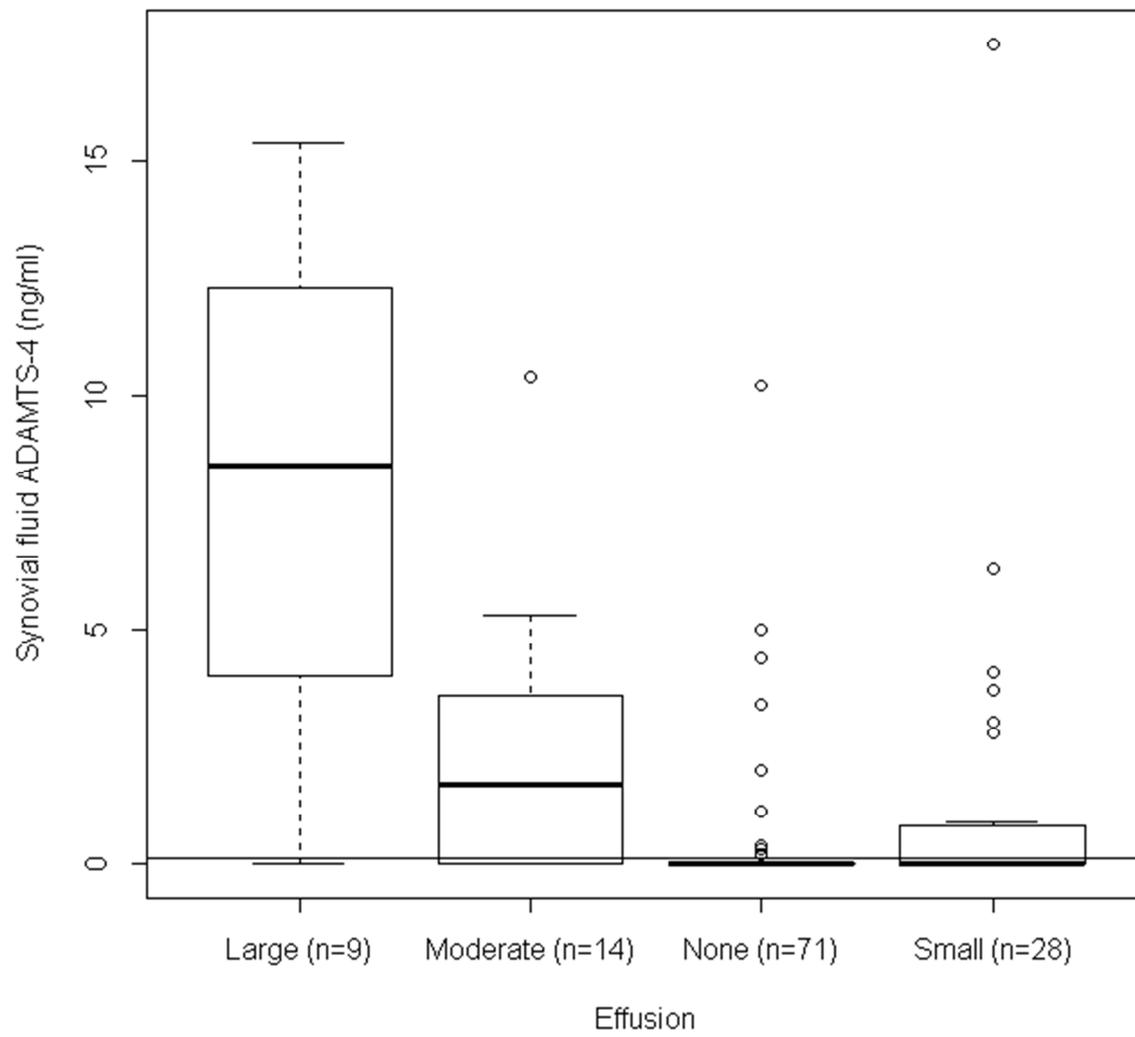


Figure 1.