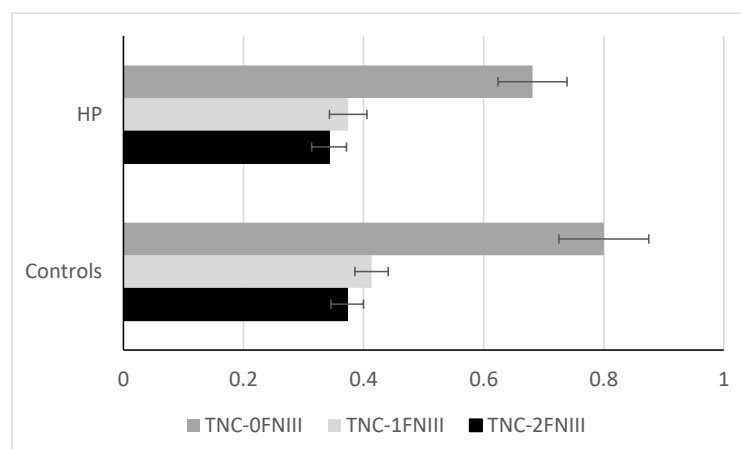
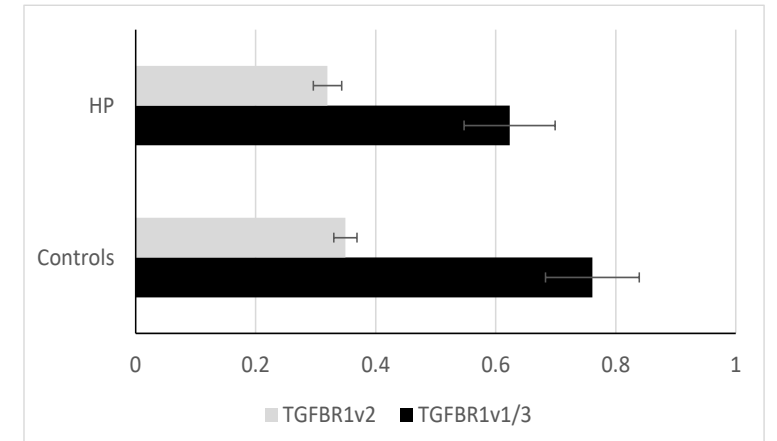
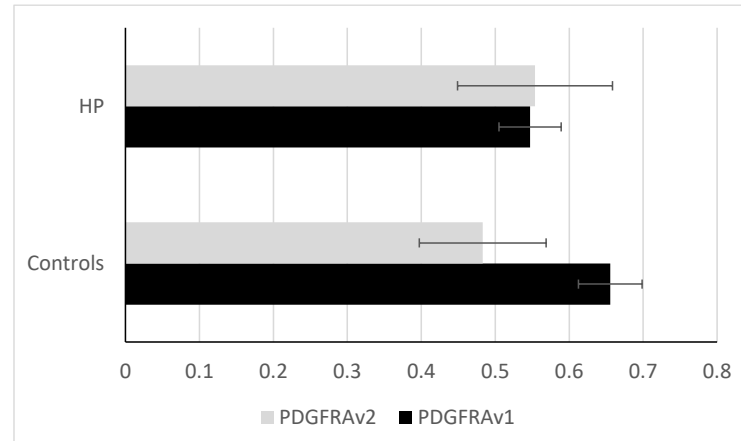
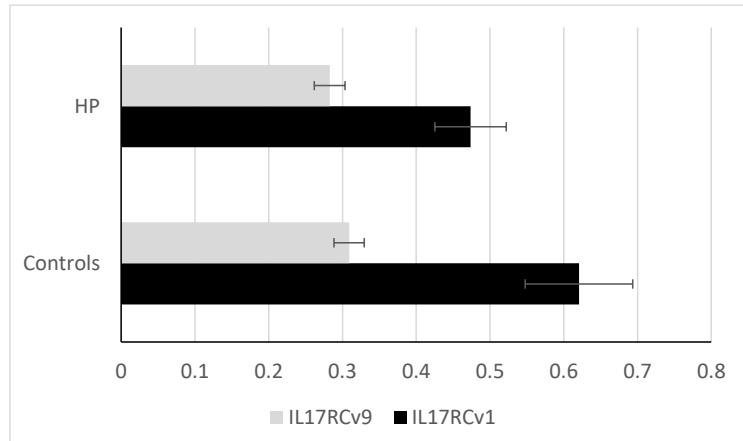
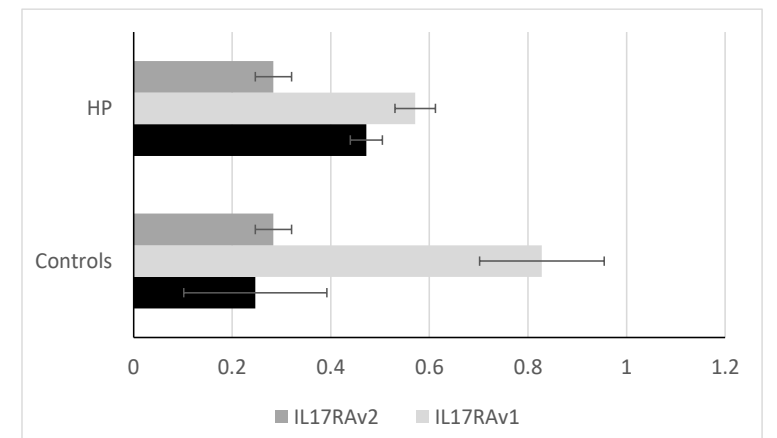
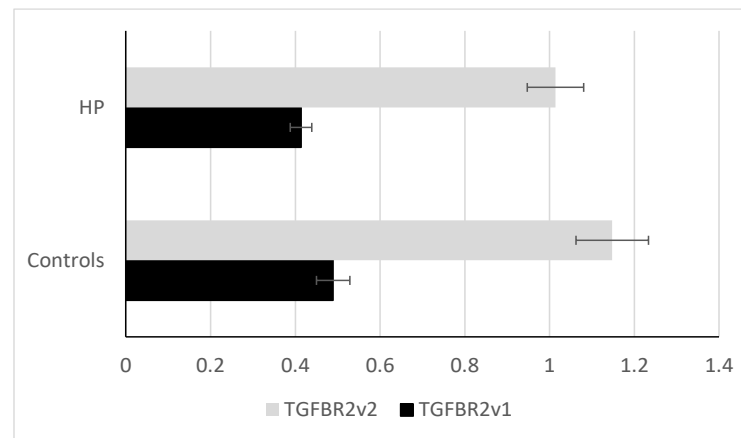
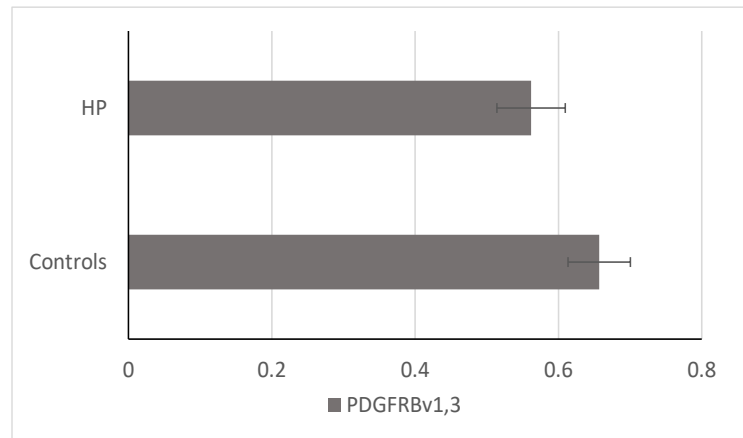
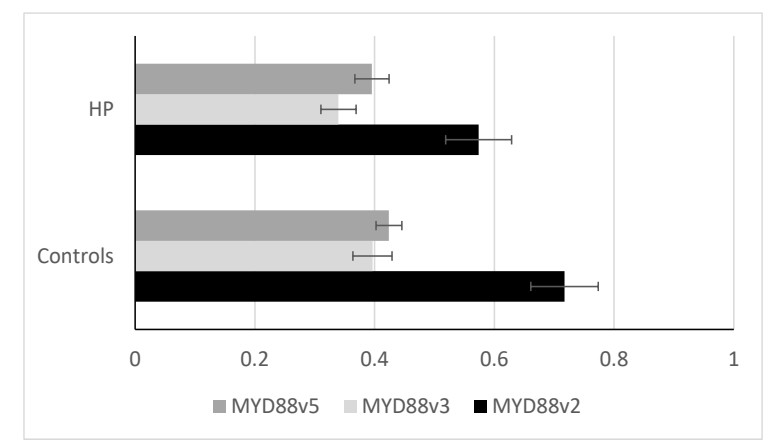
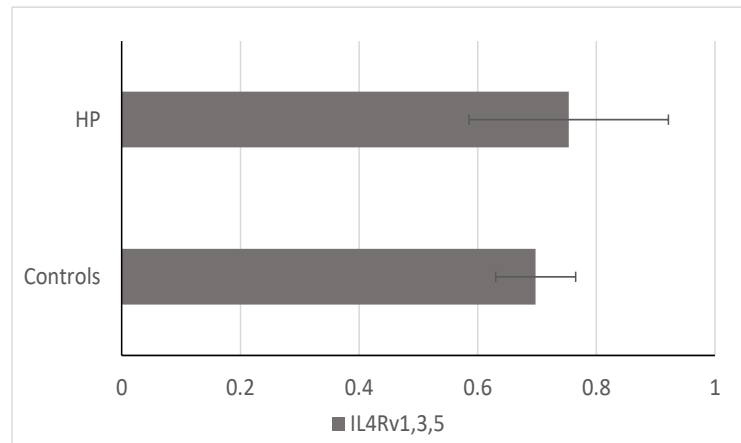
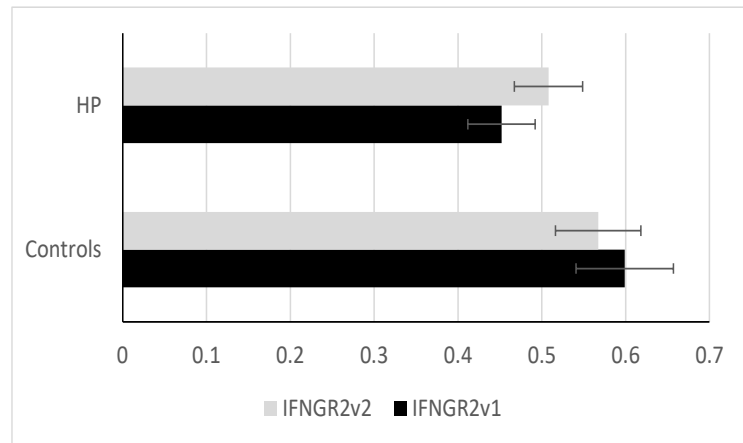
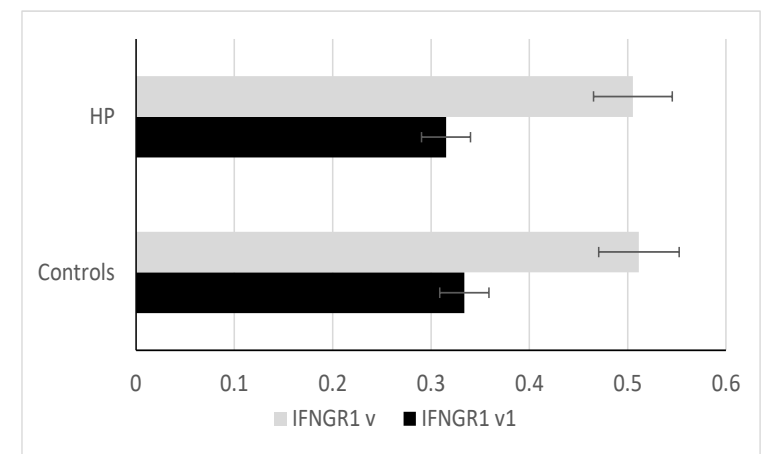
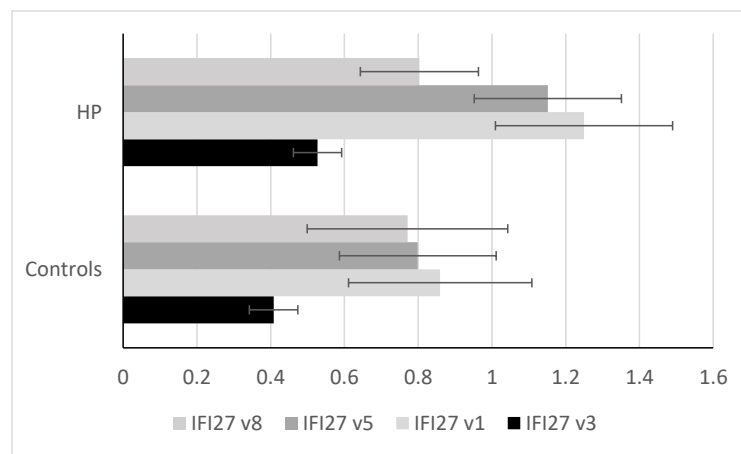
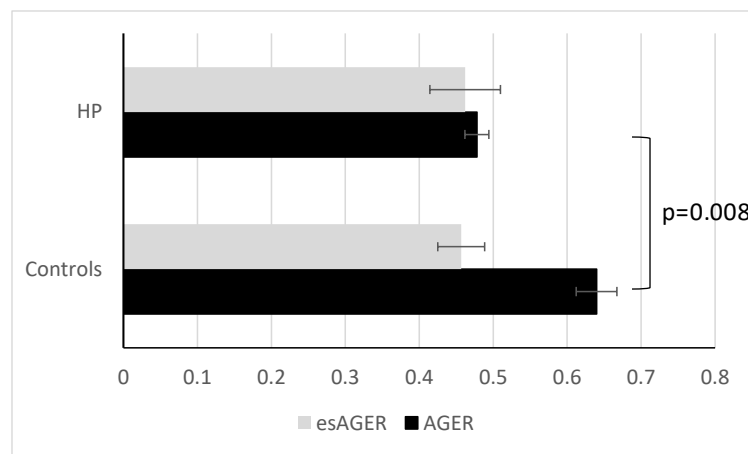


A.

B.

Gen name	Gen ID (Genbank)	Amplicon size (bp)	Samples							
			Control 1	Control 2	Control 3	Controls 4	HP1	HP2	HP3	HP4
IFI27	3429	460								
		406								
		361								
		236								
IFNGR1	3459	1457								
		1,100								
IFNGR2	3460	490								
		125								
IL17RA	23765	265								
		173								
IL17RC	84818	345								
		270								
IL4R	3566	622								
MyD88	4615	366								
		231								
		50								
PDGFRA	5156	420								
		397								
PDGFRB	5159	915								
AGER	177	183								
		139								
TGFB1	7046	762 and 773								
		528								
TGFB2	7048	379								
		284								
TNC	3371	643								
		370								
		97								
HPRT1	3251	180								

Color code	% relative to HPRT1
	0
	0.01-0.199
	0.2-0.399
	0.4-0.599
	0.6-0.799
	0.8-0.999
	1-1.199
	1.2-1.399

Figure S2. Densitometry of alternatively expressed transcripts in HP and control fibroblasts. (A) Isoform expression for each tested gene comparing control and HP samples. Error bars are the standard error. (B) Summary of the densitometric data for all genes, including the GenID and PCR product sizes. Due to the low signal, sample C3 of AGER was not included in the quantification (A). Statistical differences were assessed with the Mann-Whitney test.