Gene:							
henotype (ISSVA): Iso applies for ISSVA phe	notypes:						
. Genetic evidence .1. Case-level data .1.1. Variant evidence	посурса.						
Reference	Variant		Reported as	Mosaic Variant	Upgrade	Charterine 0.1	Score given
MID, case ID; internal data for internal data, refer to internal case-level data form	cDNA	protein	(specify phenotype designation according to reference, if applicable)	Basic Score: 0.5	Pathogenicity: 0-1.5 Prediction weak: +0; Prediction strong: +0.5; Functional effect: +0.5-1; Known LPATH: +1; Known PATH: +1.5	Clustering: 0-1 Second observation with same or other variant at same codon: +1; Neighbouring codons (+/-2): +0.5	
				0 0 0 0 0 0			0 0 0 0 0 0 0 0
				0			0
	Total score (max	kimum score for ge	enetic case level ev	idence: 12)			0
. Experimental evidence core only experimental evidence ype of vascular anomaly)	that applies for t			evidence was not or only xperimental evidence ofte		ascular anomalies, unless the	respective experiment i
ategory	Evidence Type			Default Score (Range)	Description	Reference (PMID)	Score given
2.1. Function	Biochemical function			0.5 (0-2)			
	Protein interaction			0.5 (0-2)			
	Expression			0.5 (0-2)			
2.2. Functional Alteration	Patient cells			1 (0-2)			
	Non-patient cells			0.5 (0-1)			
3. Models	Non-human model organism			2 (0-4)			
	Cell culture model			1 (0-2)			
2.4. Rescue	Rescue in human			2 (0-4)			
	Rescue in non-human model organism			2 (0-4)			
	Rescue in cell culture model			1 (0-2)			
	Rescue in patier	nt cells		1 (0-2)			
Fotal score 2.1. (maximum score fotal score 2.2. (maximum score fotal score 2.3. & 2.4. (maximum	or category: 2)	for categories: 4)					0 0 0
3. Clinical Validity Summar	v Matrix						Summary
	Genetic evidence						0
		m of genetic & exp	convincing evide Indicate PMID (Indicate PMID (nce over time (>3 yrs) (Y/N replication 1) replication 2)	1)		0
	Calculated class		Indicate PMID (replication 3)			NO EVIDENCE
		tory evidence (Y/N	1)				

3. Clinical Validity Summary Matrix					
Genetic evidence (0 to 12)	0				
Experimental evidence (0 to 6)	0				
Total points (sum of genetic & experimental evidence)	0				
Replication over time: >2 publ. w/ convincing evidence over time (>3 yrs) (Y/N)					
Indicate PMID (replication 1)					
Indicate PMID (replication 2)					
Indicate PMID (replication 3)					
Calculated classification	NO EVIDENCE				
Valid contradictory evidence (Y/N))					
if Y, list PMIDs for contradictory evidence					
if Y, describe contradictory evidence (if applicable)					
Curator classification					
Final classification (to be entered manually after panel discussion)					

Commentary:

Please enter comments regarding this gene—disease association here if applicable. This is particularly important if the curation is considered to be applicable also for other ISSVA phenotypes in addition to one that is primarily the subject of curation in part 1 ("Genetic evidence").