S5 Table. GO Terms enrichment of the 18 important, non-amyloidogenic hubs and bottlenecks, identified on the Amyloid Interactome.

	GO Term					
Protein names*	Uniprot AC	Molecular Chaperones	Autophagy	Metabolic Process	Response to Stress	Signal Transduction
14-3-3 protein zeta/delta	P63104				✓	√
Beta-arrestin-1	P49407			✓	✓	✓
Epidermal growth factor receptor	P00533			\checkmark	✓	✓
Fibronectin	P02751			\checkmark	✓	
Gamma-aminobutyric acid receptor-associated protein-like 1	Q9H0R8		✓		✓	
Gamma-aminobutyric acid receptor-associated protein-like 2	P60520		✓		✓	
Growth factor receptor-bound protein 2	P62993			✓	✓	✓
Heat shock cognate 71 kDa protein	P11142	✓	✓	\checkmark	✓	\checkmark
Hsp90 co-chaperone Cdc37	Q16543	✓	✓	✓	✓	✓
Large proline-rich protein BAG6	P46379	✓		✓	✓	✓
Mitogen-activated protein kinase 6	Q16659			✓		✓
Myc proto-oncogene protein	P01106			✓	✓	✓
Myosin-9	P35579			✓	✓	✓
NF-kappa-B essential modulator	Q9Y6K9			✓	✓	✓
Peptidyl-prolyl cis-trans isomerase NIMA-interacting 1	Q13526			✓		
Serum albumin	P02768			✓	✓	\checkmark
Transcription factor AP-1	P05412			✓	✓	✓
Vimentin	P08670					✓

GO, Gene Ontology; AC, Accession; *Protein Names were extracted from UniProt