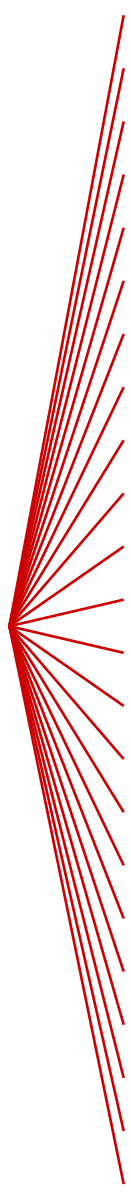


PDZ domain binding, GO:0030165



- sodium ion transmembrane transport, GO:0035725
- brush border, GO:0005903
- sodium ion transport, GO:0006814
- establishment of protein localization, GO:0045184
- potassium ion transmembrane transport, GO:0071805
- sodium channel regulator activity, GO:0017080
- phosphatidylinositol-3, GO:0043325
- frizzled binding, GO:0005109
- Ral GTPase binding, GO:0017160
- symporter activity, GO:0015293
- ventricular cardiac muscle cell action potential, GO:0086005
- positive regulation of dendrite extension, GO:1903861
- stereocilium tip, GO:0032426
- regulation of intracellular pH, GO:0051453
- organic anion transport, GO:0015711
- beta-catenin destruction complex disassembly, GO:1904886
- striated muscle contraction, GO:0006941
- epinephrine binding, GO:0051379
- postsynaptic membrane assembly, GO:0097104
- gap junction channel activity, GO:0005243
- slit diaphragm, GO:0036057
- neuropeptide signaling pathway, GO:0007218
- regulation of pH, GO:0006885