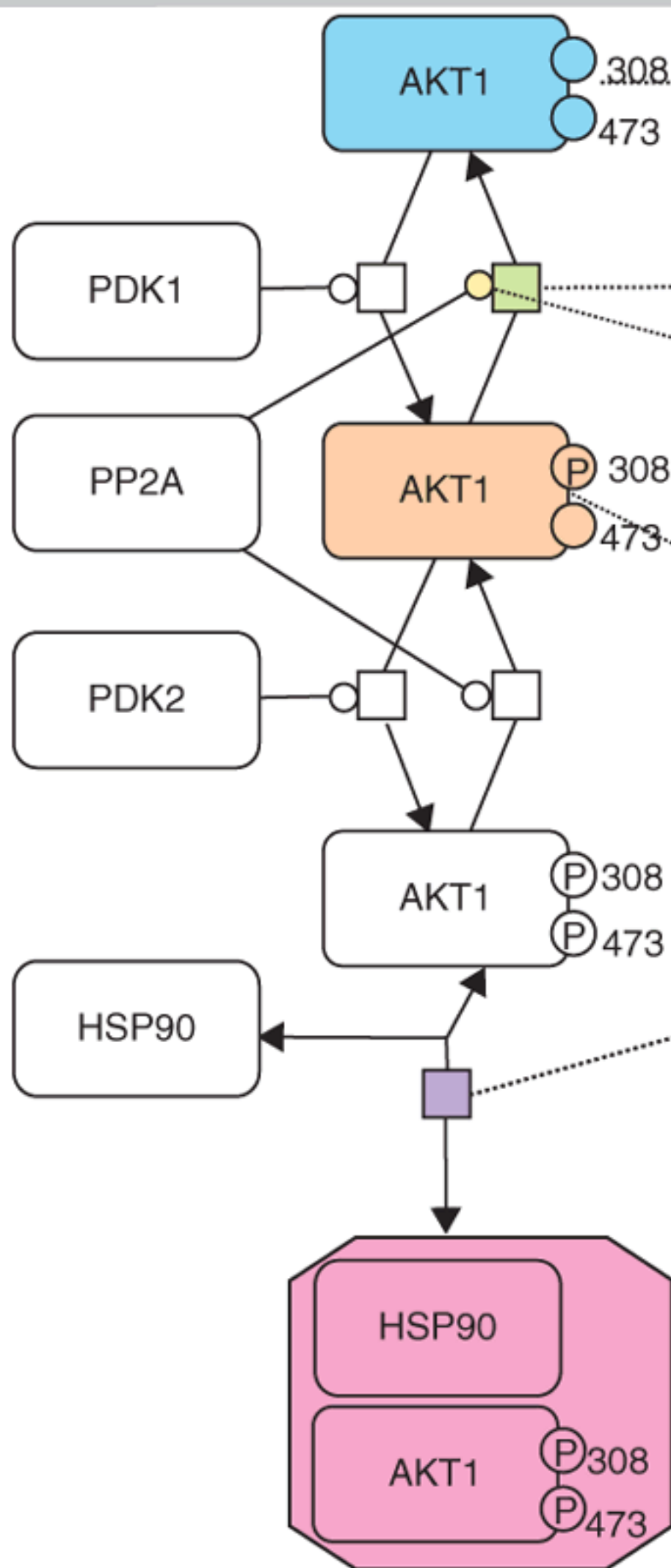


rAKT1 is a *ProteinReference*
has *standard-name* "AKT1"
has *name* "PKB"
has *xref* Uniprot-P31749

p@308 is a *ModificationFeature*
has *featureLocation* AKT1-308
has *modificationType*
phosphorylation



AKT1.1 is a *Protein*
has *proteinReference* rAKT1
has *notFeature* p@308
has *notFeature* p@473

reaction1 is a *BiochemicalReaction*
has *left* AKT1.2
has *right* AKT1.1
is *left-to-right*.

catalysis1 is a *Catalysis*
has *controller* PP2A.1
has *controlled* reaction1
has *direction* irr-left-to-right

AKT1.2 is a *Protein*
has *proteinReference* rAKT1
has *feature* p@308
has *notFeature* p@473

assembly1 is a *ComplexAssembly*
has *left* HSP90.1
has *left* AKT1.3
has *right* complex1
is *reversible*

complex1 is a *Complex*
has *component* AKT1.4
has *component* HSP90.2

HSP90.2 is a *Protein*
has *proteinReference* rHSP90
is *boundTo* AKT1.4

AKT1.4 is a *Protein*
has *proteinReference* rAKT1
has *feature* p@308
has *feature* p@473
is *boundTo* HSP90.2