

# MET phosphorylates SHC1-2

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## Introduction

Reactome is open-source, open access, manually curated and peer-reviewed pathway database. Pathway annotations are authored by expert biologists, in collaboration with Reactome editorial staff and cross-referenced to many bioinformatics databases. A system of evidence tracking ensures that all assertions are backed up by the primary literature. Reactome is used by clinicians, geneticists, genomics researchers, and molecular biologists to interpret the results of high-throughput experimental studies, by bioinformaticians seeking to develop novel algorithms for mining knowledge from genomic studies, and by systems biologists building predictive models of normal and disease variant pathways.

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## Literature references

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Reactome database release: 70

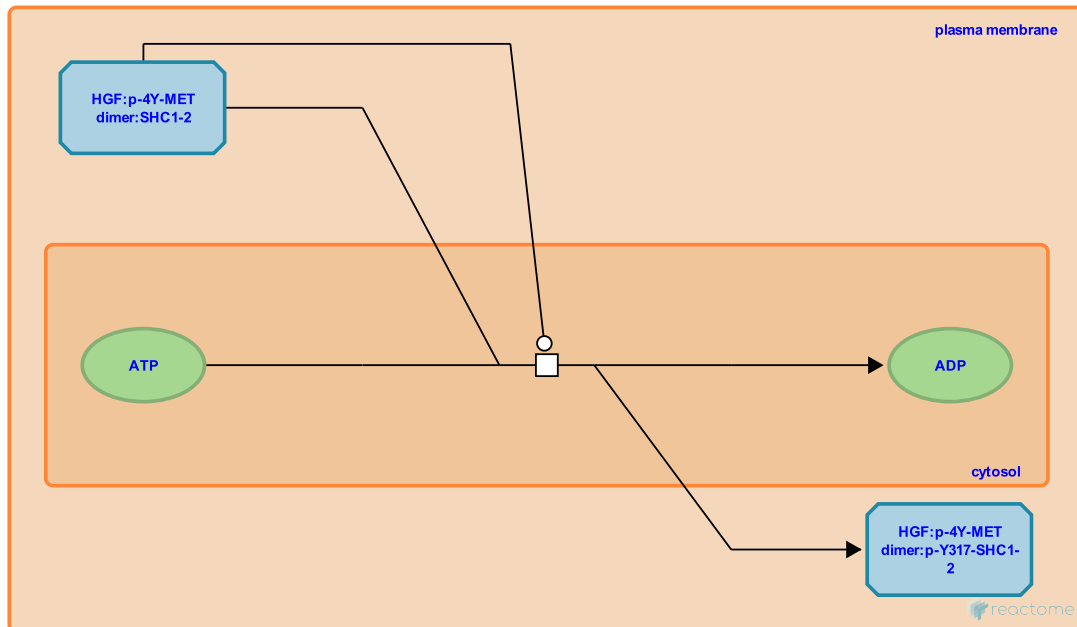
This document contains 1 reaction ([see Table of Contents](#))

## MET phosphorylates SHC1-2 [↗](#)

**Stable identifier:** R-HSA-8851890

**Type:** transition

**Compartments:** cytosol, plasma membrane



Activated MET receptor phosphorylates the splicing isoform 2 of the adapter protein SHC1 (SHC1-2) on tyrosine residue Y317 (Pelicci et al. 1995).

### Literature references

Pelicci, G., Giordano, S., Zhen, Z., Salcini, AE., Lanfrancone, L., Bardelli, A. et al. (1995). The mitogenic and mitogenic responses to HGF are amplified by the Shc adaptor protein. *Oncogene*, 10, 1631-8. [↗](#)

### Editions

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