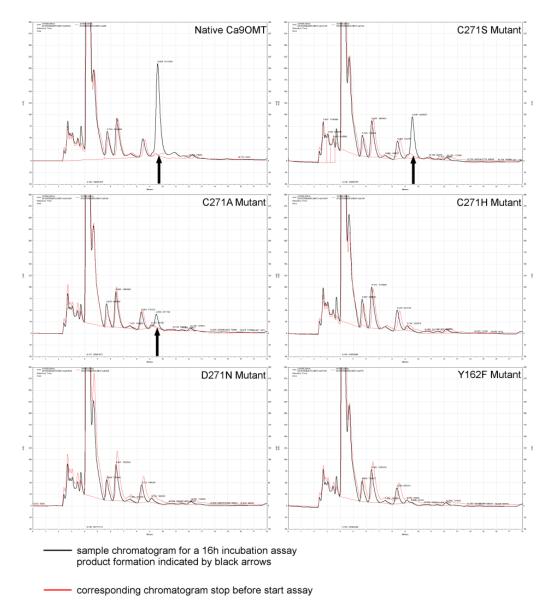
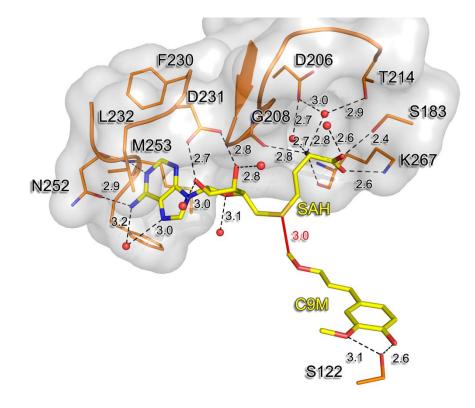
## **Supplementary Material**



Supplementary Figure S1. Representative chromatograms.



Supplementary Figure S2. SAM binding mode.

## Supplementary Table S1. SAM interactions.

## Residues complexing the purine ring system of SAM/SAH

Y162: non-polar cavity

L232: non-polar cavity

N252: H-bond between side chain and purine ring amino function(N6)2.9 Å

M253: non-polar cavity

## Residues complexing the ribose ring of SAM/SAH

**<u>D231:</u>** two direct H-bond interactions between side chain and ribose hydroxyl functions 2.8 Å and 2.7 Å

**<u>S176</u>** bridged H-bond via HOH586A, distance S176 HOH586A: 3.0 Å distance HOH586A and hydroxyl function of SAM(O3): 2.8 Å

**<u>H234</u>**: bridged H-bond via HOH586A distance H234 HOH586A: 3.1 Å distance HOH586A and hydroxyl function of SAM(O2): 3.0 Å

**D272:** bridged H-bond via HOH568A distance D272 HOH568A: 2.9 Å distance HOH568A and ring bound oxygen(O4) of SAM: 3.1 Å

**W273:** bridged H-bond via HOH568A distance W273 HOH568A: 2.9 Å distance HOH568A and ring bound oxygen(O4) of SAM: 3.1 Å

Residues complexing the thioether function of SAM

V205-T214: several non-polar interactions

Residues complexing the methionine carboxyl and amino functions

**<u>S183:</u>** H-bond between side chain and SAM carboxyl function (O) 2.4 Å

G208: H-bond between backbone oxygen and SAM amino function (N)\_2.8 Å

<u>**K267:**</u> H-bond between side chain and SAM carboxyl function(O): 2.6 Å\_H-bond between backbone oxygen and SAM amino function 2.7 Å

**Supplementary Table S2:** Results of isothermal titration calorimetry measurements (ITC) all values derived from three independent measurements. Error ranges indicate the estimated standard deviations from triplicate measurements.

Combination of Ligand and Protein	N [sites]	К <sub>ь</sub> [М⁻¹]	К <sub>d</sub> [µМ]	∆H⁰ [kj/mol]	-T∆S⁰ [kj/mol/ deg]
A:experimental setup 1 sample cell: Ca9OMT syringe:	0.779 ±0.212	1.95•10⁵ ±0.38•10⁵	5.26 ±1.02	-43.7 ±5.7	13.0 ±5.3
coniferyl alcohol <u>B:experimental</u> <u>setup 2</u> sample cell: Ca9OMT+coniferyl alcohol syringe: SAM	0.525 ±0.057	2.45•10 <sup>6</sup> ±0.85•10 <sup>6</sup>	0.45 ±0.16	-55.1 ±3.5	18.2 ±4.4
C:experimental setup 3 sample cell: Ca9OMT syringe: SAM	n.d.	n.d.	n.d.	n.d.	n.d.
D:experimental setup 4 sample cell: Ca9OMT+SAM syringe: coniferyl alcohol	0.865 ±0.046	3.96•10 <sup>5</sup> ±1.95•10 <sup>5</sup>	2.94 ±1.29	-62.2 ±4.0	29.9 ±5.3