## 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 $\AA$ |
| M253: non-polar cavity |
| Residues complexing the ribose ring of SAM/SAH |
| D231: two direct H-bond interactions between side chain and ribose hydroxyl functions $2.8 \AA$ and $2.7 \AA$ |
| S176: bridged H-bond via HOH586A,distance S176 HOH586A: 3.0 Å_distance HOH586A and hydroxyl function of SAM(O3): $2.8 \AA$ |
| H234: bridged H-bond via HOH586A distance H234 HOH586A: 3.1 Å distance HOH586A and hydroxyl function of SAM(O2): $3.0 \AA$ |
| D272: bridged H-bond via HOH568A distance D272 HOH568A: 2.9 Å distance HOH568A and ring bound oxygen(O4) of SAM: $3.1 \AA$ |
| W273: bridged H-bond via HOH568A distance W273 HOH568A: 2.9 Å distance HOH568A and ring bound oxygen(O4) of SAM: $3.1 \AA$ |
| Residues complexing the thioether function of SAM |
| V205-T214: several non-polar interactions |
| Residues complexing the methionine carboxyl and amino functions |
| S183: H-bond between side chain and SAM carboxyl function (O)_2.4 $\AA$ |
| G208: H-bond between backbone oxygen and SAM amino function (N)_2.8 $\AA$ |
| K267: H-bond between side chain and SAM carboxyl function(O): $2.6 \AA$ _H-bond between backbone oxygen and SAM amino function $2.7 \AA$ |

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] | $\begin{aligned} & \mathbf{K}_{\mathrm{b}} \\ & {\left[\mathbf{M}^{-1}\right]} \end{aligned}$ | $K_{d}$ <br> [ $\mu \mathrm{M}$ ] | $\Delta \mathbf{H}^{0}$ [kj/mol] | $\begin{aligned} & -\mathrm{T} \Delta \mathbf{S}^{0} \\ & {[\mathrm{kj} / \mathrm{mol} /} \\ & \mathrm{deg}] \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: |
| A:experimental setup 1 sample cell: Ca9OMT syringe: coniferyl alcohol | $\begin{aligned} & 0.779 \\ & \pm 0.212 \end{aligned}$ | $\begin{aligned} & 1.95 \cdot 10^{5} \\ & \pm 0.38 \cdot 10^{5} \end{aligned}$ | $\begin{aligned} & 5.26 \\ & \pm 1.02 \end{aligned}$ | $\begin{aligned} & -43.7 \\ & \pm 5.7 \end{aligned}$ | $\begin{aligned} & 13.0 \\ & \pm 5.3 \end{aligned}$ |
| $\begin{aligned} & \hline \text { B:experimental } \\ & \hline \text { setup 2 } \\ & \text { sample cell: } \\ & \text { Ca9OMT+coniferyl } \\ & \text { alcohol } \\ & \text { syringe: } \\ & \text { SAM } \end{aligned}$ | $\begin{aligned} & 0.525 \\ & \pm 0.057 \end{aligned}$ | $\begin{array}{l\|} \hline 2.45 \cdot 10^{6} \\ \pm 0.85 \cdot 10^{6} \end{array}$ | $\begin{aligned} & 0.45 \\ & \pm 0.16 \end{aligned}$ | $\begin{aligned} & -55.1 \\ & \pm 3.5 \end{aligned}$ | $\begin{aligned} & 18.2 \\ & \pm 4.4 \end{aligned}$ |
| $\begin{aligned} & \hline \text { C:experimental } \\ & \hline \text { setup } 3 \\ & \text { sample cell: } \\ & \text { Ca9OMT } \\ & \text { syringe: } \\ & \text { SAM } \\ & \hline \end{aligned}$ | n.d. | n.d. | n.d. | n.d. | n.d. |
| $\begin{aligned} & \hline \text { D:experimental } \\ & \hline \text { setup 4 } \\ & \text { sample cell: } \\ & \text { Ca9OMT }+ \text { SAM } \\ & \text { syringe: } \\ & \text { coniferyl alcohol } \\ & \hline \end{aligned}$ | $\begin{aligned} & 0.865 \\ & \pm 0.046 \end{aligned}$ | $\begin{aligned} & 3.96 \cdot 10^{5} \\ & \pm 1.95 \cdot 10^{5} \end{aligned}$ | $\begin{aligned} & 2.94 \\ & \pm 1.29 \end{aligned}$ | $\begin{aligned} & -62.2 \\ & \pm 4.0 \end{aligned}$ | $\begin{aligned} & 29.9 \\ & \pm 5.3 \end{aligned}$ |

