Supplementary Figures

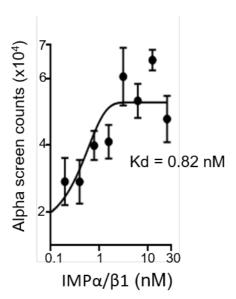


Fig. S1. High affinity recognition of DENV NS5 by the IMP α / β 1 heterodimer. AlphaScreen technology was used to determine the dissociation constant (Kd) of binding of IMP α / β 1 (prebound IMP α / β 1 heterodimer with biotinylated IMP β 1) to His6-DENV NS5 (30 nM). Data represent the mean +/- SD for triplicate wells from a single typical experiment, from a series of 2 independent experiments (see Table 1 column 1 for pooled data).

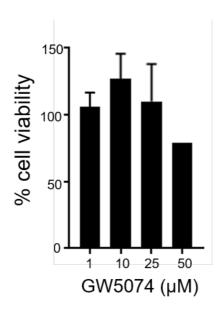


Fig. S2. Lack of toxicity of GW5074 at concentrations effective in inhibiting flavivirus. Cell viability was determined by addition of XTT reagent (Sigma-Aldrich) following compound treatments, as indicated. Cell survival is plotted relative to an untreated control. Data represent the mean +/- SD for duplicate wells from a single typical experiment, from a series of 2 independent experiments.

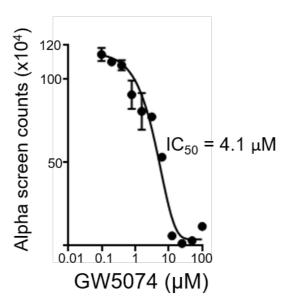


Fig. S3. GW5074 inhibits recognition of SV40 T-antigen by the IMP $\alpha/\beta1$ heterodimer. AlphaScreen technology was used to determine the IC50 for inhibition by GW5074 of T-ag binding to IMP $\alpha/\beta1$. Data represent the mean +/- SD for triplicate wells from a single typical experiment, from a series of 2 independent experiments.

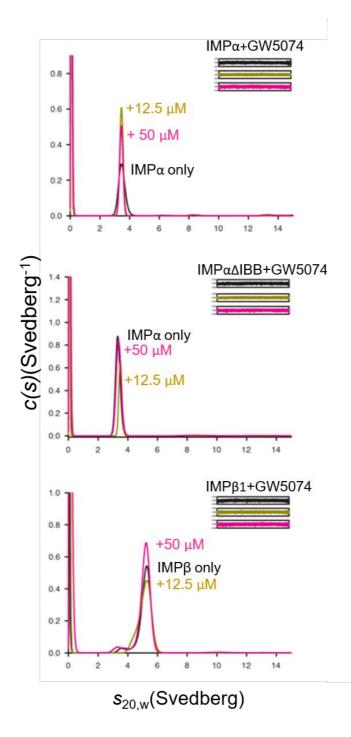


Fig. S4. GW5074 does not alter the sedimentation coefficients of IMP α , IMP $\alpha\Delta$ IBB or IMP β 1. Sedimentation velocity analytical ultracentrifugation experiments were performed on purified recombinant IMP α , IMP $\alpha\Delta$ IBB and IMP β 1, in the absence or presence of GW5074. The continuous sedimentation coefficient distribution [(c)s] was plotted as a function of s20,w for IMPs alone, in the presence of the indicated concentrations of GW5074. The residual plots are shown in insets. Results are from a single typical experiment, from a series of 2 independent experiments.

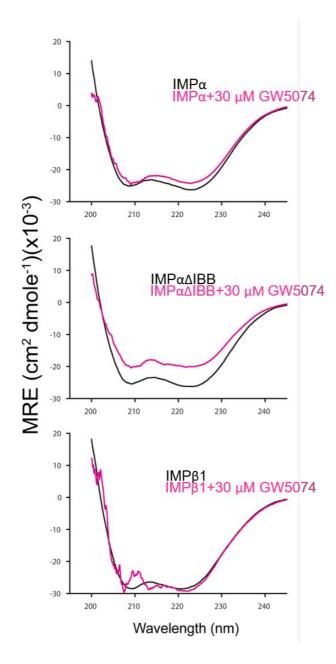


Fig. S5. CD spectra for IMPs in the absence and presence of GW5074. CD spectra were collected for IMP α , IMP $\alpha\Delta$ IBB and IMP β 1 in the absence or presence of 30 μ M GW5074. The plots are representative of 2 independent experiments.

Supplementary Materials

Table S1. Hydrodynamic properties of recombinant IMP proteins.

Protein	Parameter			
	$M_{ m r}^{ m a}$	M^{b}	S20,w ^c	f/f_0 d
ΙΜΡα	58072	50316	3.5	2.4
ІМРβ1	98557	106358	5.3	2.1
ΙΜΡα/β1	156629	155055	6.7	2.7

^a Relative molecular weight calculated from the amino acid sequence.

^b Molar mass determined from the ordinate maximum of c(M) distribution best fits (data not shown).

^c Standardized sedimentation coefficient taken from the ordinate maximum of the c(s) distribution best fits (Fig. 2C).

 $^{^{}m d}$ Frictional coefficient calculated from $s_{20, w}$ using the \overline{v} method employing SEDNTERP.