

Supplemental materials for “Preclinical Characterization of BMS-791325, an Allosteric Inhibitor of Hepatitis C Virus NS5B Polymerase” (Lemm J, *et al.*)

Contents

| | |
|---|---|
| Table S1. Triple Combination Using BMS-791325 ('325) With IFN α and Ribavirin (RBV)..... | 2 |
| Table S2. Combination of BMS-791325 ('325) With the NS3 Protease Inhibitor Asunaprevir (ASV)..... | 3 |
| Table S3. Combination of BMS-791325 ('325) With the NS5A Inhibitor Daclatasvir (DCV)..... | 4 |
| Table S4. Combination of BMS-791325 ('325) With the Palm Site 2 NS5B Inhibitor HCV-796..... | 5 |
| Table S5. Combination of BMS-791325 ('325) With the NS5B Nucleoside Inhibitor NM-283..... | 6 |
| Table S6. Triple Combination Using BMS-791325 ('325) With the NS5A inhibitor Daclatasvir (DCV) and the NS3 Protease Inhibitor Asunaprevir (ASV) | 7 |
| Database Consensus Sequences for GT 2b, 3a, 4a, 5a and 6a NS5B Proteins | 8 |

Table S1. Triple Combination Using BMS-791325 ('325) With IFN α and Ribavirin (RBV)

| Expt | '325 EC ₅₀ , nM | IFN α EC ₅₀ , U/mL | RBV EC ₅₀ , nM | Combination Indices (confidence interval) | | | Overall Result |
|------|-------------------------------|---|------------------------------|--|----------------------|----------------------|------------------------|
| | | | | 50% effective | 75% effective | 90% effective | |
| 1 | 4.4 | 1.6 | 24566 | 0.97 (0.85, 1.09) | 0.97 (0.80, 1.14) | 1.02 (0.74, 1.29) | Additivity |
| 2 | 4.3 | 7.1 | 45323 | 1.04 (0.95, 1.13) | 0.95 (0.84, 1.06) | 0.88 (0.73, 1.04) | Additivity |
| 3 | 4.1 | 9.1 | 53362 | 0.93 (0.84, 1.01) | 0.82 (0.72, 0.93) | 0.76 (0.61, 0.91) | Additivity/ Synergy |

Table S2. Combination of BMS-791325 ('325) With the NS3 Protease Inhibitor Asunaprevir (ASV)

| Expt | '325 EC ₅₀ , uM | ASV EC ₅₀ , nM | ASV:'325 | Combination Indices (confidence interval) | | | Overall Result |
|------|-------------------------------|------------------------------|----------|--|----------------------|----------------------|----------------|
| | | | | 50% effective | 75% effective | 90% effective | |
| 1 | 0.003 | 0.7 | 1:1 | 1.05 (0.98, 1.12) | 0.96 (0.86, 1.06) | 0.88 (0.74, 1.02) | Additivity |
| | | | 2.5:1 | 0.95 (0.87, 1.03) | 0.99 (0.88, 1.11) | 1.04 (0.84, 1.23) | Additivity |
| | | | 1:2.5 | 0.98 (0.91, 1.06) | 0.99 (0.88, 1.09) | 0.99 (0.82, 1.15) | Additivity |
| 2 | 0.003 | 0.7 | 1:1 | 1.04 (0.96, 1.12) | 0.95 (0.84, 1.05) | 0.86 (0.71, 1.01) | Additivity |
| | | | 2.5:1 | 0.97 (0.89, 1.05) | 1.00 (0.88, 1.12) | 1.03 (0.84, 1.22) | Additivity |
| | | | 1:2.5 | 0.97 (0.89, 1.05) | 0.98 (0.87, 1.10) | 1.00 (0.82, 1.18) | Additivity |
| 3 | 0.004 | 0.9 | 1:1 | 1.00 (0.92, 1.07) | 0.96 (0.86, 1.06) | 0.92 (0.77, 1.08) | Additivity |
| | | | 2.5:1 | 1.07 (1.00, 1.14) | 1.00 (0.91, 1.09) | 0.93 (0.80, 1.07) | Additivity |
| | | | 1:2.5 | 1.02 (0.93, 1.11) | 1.03 (0.90, 1.16) | 1.04 (0.84, 1.24) | Additivity |

Table S3. Combination of BMS-791325 ('325) With the NS5A Inhibitor Daclatasvir (DCV)

| Expt | DCV EC ₅₀ , nM | '325 EC ₅₀ , uM | '325:DCV | Combination Indices (confidence interval) | | | Overall Result |
|------|------------------------------|-------------------------------|----------|--|----------------------|----------------------|-------------------------|
| | | | | 50% effective | 75% effective | 90% effective | |
| 1 | 0.010 | 0.015 | 1:1 | 0.88 (0.81, 0.95) | 0.83 (0.73, 0.93) | 0.79 (0.64, 0.94) | Synergy |
| | | | 1:2.5 | 0.80 (0.72, 0.88) | 0.79 (0.68, 0.89) | 0.78 (0.62, 0.94) | Synergy |
| | | | 2.5:1 | 0.97 (0.90, 1.04) | 0.97 (0.87, 1.07) | 0.96 (0.80, 1.13) | Additivity |
| 2 | 0.007 | 0.013 | 1:1 | 0.81 (0.76, 0.86) | 0.88 (0.80, 0.95) | 0.95 (0.82, 1.08) | Synergy / Additivity |
| | | | 1:2.5 | 0.66 (0.62, 0.70) | 0.65 (0.59, 0.72) | 0.65 (0.55, 0.75) | Synergy |
| | | | 2.5:1 | 0.95 (0.88, 1.01) | 0.98 (0.90, 1.07) | 1.02 (0.88, 1.16) | Additivity |
| 3 | 0.006 | 0.010 | 1:1 | 0.85 (0.77, 0.93) | 0.88 (0.77, 0.98) | 0.90 (0.73, 1.08) | Synergy / Additivity |
| | | | 1:2.5 | 0.84 (0.79, 0.90) | 0.81 (0.72, 0.91) | 0.79 (0.64, 0.93) | Synergy |
| | | | 2.5:1 | 0.92 (0.85, 0.99) | 0.92 (0.82, 1.01) | 0.92 (0.77, 1.06) | Synergy / Additivity |

Table S4. Combination of BMS-791325 ('325) With the Palm Site 2 NS5B Inhibitor HCV-796

| Expt | '325 EC ₅₀ , uM | HCV-796 EC ₅₀ , uM | HCV- 796:'325 | Combination Indices (confidence interval) | | | Overall Result |
|------|-------------------------------|----------------------------------|------------------|--|----------------------|----------------------|----------------|
| | | | | 50% effective | 75% effective | 90% effective | |
| 1 | 0.005 | 0.013 | 1:1 | 0.81 (0.77, 0.85) | 0.76 (0.70, 0.81) | 0.70 (0.63, 0.78) | Synergy |
| | | | 2.5:1 | 0.82 (0.78, 0.87) | 0.79 (0.73, 0.84) | 0.75 (0.66, 0.83) | Synergy |
| | | | 1:2.5 | 0.88 (0.84, 0.91) | 0.78 (0.73, 0.84) | 0.70 (0.63, 0.78) | Synergy |
| 2 | 0.004 | 0.008 | 1:1 | 0.79 (0.75, 0.83) | 0.66 (0.61, 0.71) | 0.55 (0.48, 0.62) | Synergy |
| | | | 2.5:1 | 0.79 (0.75, 0.84) | 0.70 (0.65, 0.76) | 0.62 (0.54, 0.70) | Synergy |
| | | | 1:2.5 | 0.76 (0.72, 0.81) | 0.69 (0.64, 0.75) | 0.63 (0.55, 0.71) | Synergy |
| 3 | 0.003 | 0.011 | 1:1 | 0.83 (0.76, 0.90) | 0.73 (0.64, 0.82) | 0.64 (0.52, 0.77) | Synergy |
| | | | 2.5:1 | 0.74 (0.68, 0.80) | 0.71 (0.63, 0.80) | 0.69 (0.57, 0.82) | Synergy |
| | | | 1:2.5 | 0.88 (0.80, 0.95) | 0.84 (0.74, 0.93) | 0.80 (0.66, 0.94) | Synergy |

Table S5. Combination of BMS-791325 ('325) With the NS5B Nucleoside Inhibitor NM-283

| Expt | '325 EC ₅₀ , uM | NM-283 EC ₅₀ , uM | NM- 283:'325 | Combination Indices (confidence interval) | | | Overall Result |
|------|-------------------------------|---------------------------------|-----------------|--|----------------------|----------------------|-------------------------|
| | | | | 50% effective | 75% effective | 90% effective | |
| 1 | 0.003 | 1.2 | 1:1 | 0.95 (0.85, 1.05) | 0.88 (0.75, 1.01) | 0.81 (0.63, 1.00) | Additivity / Synergy |
| | | | 2.5:1 | 0.77 (0.68, 0.86) | 0.84 (0.70, 0.98) | 0.92 (0.68, 1.15) | Synergy / Additivity |
| | | | 1:2.5 | 0.80 (0.71, 0.89) | 0.82 (0.69, 0.95) | 0.85 (0.64, 1.06) | Synergy / Additivity |
| 2 | 0.003 | 1.3 | 1:1 | 0.53 (0.47, 0.59) | 0.59 (0.50, 0.69) | 0.67 (0.50, 0.83) | Synergy |
| | | | 2.5:1 | 0.44 (0.39, 0.50) | 0.53 (0.44, 0.62) | 0.64 (0.47, 0.80) | Synergy |
| | | | 1:2.5 | 0.43 (0.37, 0.49) | 0.59 (0.47, 0.70) | 0.80 (0.56, 1.04) | Synergy / Additivity |
| 3 | 0.003 | 1.0 | 1:1 | 0.76 (0.68, 0.84) | 0.76 (0.65, 0.87) | 0.75 (0.58, 0.93) | Synergy |
| | | | 2.5:1 | 0.77 (0.69, 0.85) | 0.75 (0.64, 0.86) | 0.73 (0.57, 0.90) | Synergy |
| | | | 1:2.5 | 0.60 (0.54, 0.66) | 0.63 (0.54, 0.71) | 0.66 (0.52, 0.80) | Synergy |

Table S6. Triple Combination Using BMS-791325 ('325) With the NS5A inhibitor Daclatasvir (DCV) and the NS3 Protease Inhibitor Asunaprevir (ASV)

| Expt | '325 EC ₅₀ , nM | DCV EC ₅₀ , nM | ASV EC ₅₀ , nM | Combination Indices (confidence interval) | | | Overall Result |
|------|-------------------------------|------------------------------|------------------------------|--|----------------------|----------------------|----------------|
| | | | | 50% effective | 75% effective | 90% effective | |
| 1 | 5.4 | 0.003 | 1.2 | 1.00 (0.92, 1.08) | 1.05 (0.93, 1.17) | 1.11 (0.91, 1.30) | Additivity |
| 2 | 2.3 | 0.002 | 0.7 | 0.93 (0.85, 1.01) | 1.01 (0.88, 1.14) | 1.10 (0.88, 1.31) | Additivity |
| 3 | 2.8 | 0.002 | 0.8 | 1.01 (0.94, 1.08) | 0.97 (0.88, 1.06) | 0.93 (0.79, 1.07) | Additivity |

'325, BMS-791325; DCV, daclatasvir; ASV, asunaprevir.

Database Consensus Sequences for GT 2b, 3a, 4a, 5a and 6a NS5B Proteins

GT-2b

MSTSYSWTGALITPCGPEEEKLPINPLSNLSLMRFHNKVYSTTSRSASQRAKKVTFDRVQVLDTHYDSVLQDVKRA
ASKVSARLLSVEEACALTPPHSAKSRYGFGAKEVRSLRRAVNHRSVWEDLLEDQHTPIDTTIMAKNEVFCVDPA
KGGKKSARLIVYPDLGVRVCEKMALYDIAQKLPKAVMGSSYGFQYSPAERVDFLLKAWGSKKDPMGFSYDTRCF
DSTVTERDIRTEESIQACSLPQEARTVIHSLTERLYVGGPMINSKGQSCGYRRCRASGVFTTSMGNTMTCYIKAL
AACKAAGIVDPIMLVCGDDLVIIVISESQGNEEDERNLRAFTEAMTRYSAPPGDLPRPEYDLELITSCSSNVSVLDP
RGRRRYYLTRDPTTISRRAAWETVRHSPVNSWLGNIQYAPTIWVRMVIMTHFFAILLAQDTLNQNLNFEMYGAVY
SVNPLDLPALIERIHGLDAFSLHTYSPHELRSVAATLRKLGAPPLRAWKSRARAVRASLIAQGGGRASVCGRYLFNW
AVRTKCLKLTPLPEASRLDLSG WFTVGAGGGDIFHSVSHARPRLLL

GT-3a

MSMSYSWTGALITPCSAEEEEKLPISPLSNLSLLRHHNLVYSTSSRSASQRQKKVTFDRLQVLDDHYKTVLQEVKER
ASRVKARTLTIEEACALVPPHSARSKFGYSAKDVRSLSSKAINQIRSVWEDLLEDTTTTPIPTTIMAKNEVFCVDPTK
GGRKPTRLIVYPDLGVRVCEKRALYDVIQKLSIETMGPAYGFQYSPQQRVERLLKMWTSKKTPLGFSYDTRCFDS
TVTEQDIRVEEIIYQCCNLEPEARKVISSLTERLYCGGPMFNSKGAQCCGYRRCRASGVLPSTFGNTITCYIKATAA
ARAAGLRNPEFLVCGDDLVIIVAESDGVDEDRATLRAFTEVMTRYSAAPGDAPQPTYDLELITSCSSNVSVARDDK
GKRYYYLTRDATTPLARAAWETARHTPVNSWLGNIIMYAPTIWVRMVMMTHFFSILQSQEILDRPLDFEMYGATYS
VTPLDLPALIERLHGLSAFTLHSPVELNRVAGTLRKLGCPLRAWRHRARAVRAKLIQGGKAKICGLYLFNWA
VRTKTKLTPLPATGQLDLSSWFTVGVGNDIYHSVSRARTRYLL

GT-4a

MSMSYSWTGALVTPCAAEEKLPISPLSNLSLLRHHNMVYATTTSAVTRQKKVTFDRLQVVDNHYNLKEIKAR
ASRVKARLLTTEEACDLTPPHSAKSKFGYGAKDVRSHSRKAINHINSVWEDLLEDNNTPIPTTIMAKNEVFAVNPA
KGGKPARLIVYPDLGVRVCEKRALHDVINQLPKAVMGAAYGFQYSPAQRVEFLLTSWKSCKTPMGFSYDTRCF
DSTVTEKDIRTEEEVYQCCDLEPEARKVIAALTERLYVGGPMHNSKGDLCGYRRCRASGVYTTSTFGNTLTCYLKA
TAAIKAAGLRDCTMLVCGDDLVIIVAESDGVVEDNRALRAFTEAMTRYSAPPGDAPQPAYDLELITSCSSNVSVAH
DATGKKVYYLTRDPETPLARAAWETVRHTPVNSWLGNIIVYAPTIWVRMVLMTTHFFSILQSQEALDFDYMVG
TYSITPLDLPALIIQRLHGLSAFTLHGYSPELNRVAGSLRKLGVPLRAWRHRARAVRAKLIQGGKAKICGIYLFN
WAVKTKLKLTPLPAAANLDLSSWFTVGAGGGDIYHSVSHARPRYLL

GT-5a

MSMSYSWTGALITPCSADEEKLPINPLSNLTLRHHNLVYSTSSRSAGLRQKKVTFDRLQVLDDHYREVVDKMKRL
ASKVKARLLPLEEACGLTPPHSARSKYGYGAKEVRSLDKKALNHKGVWQDLDLSDTPTTIMAKNEVFAVEPS
KGGKPARLIVYPDLGVRVCEKRALYDVAQKLPTALMGPSYGFQYSPAQRVEFLLKAWKSKKTPMAFSYDTRCF
DSTVTEHDIMTEESIQSCDLQPEARAAIRSLTQRLYCGGPMYNSKGQQCCGYRRCRASGVFTTSMGNTMTCYIK
ALASCRAAKLRDCTLLVCGDDLVAICESQGTHEDEASLRAFTEAMTRYSAPPGDPPVPAYDLELVTSCSSNVSV
RDASGNRVYYLTRDPQVPLARAAWETAKHSPVNSWLGNIIMYAPTLWARIVLMTTHFFSVLQSQEQLKALAFEMY
GSVYSVTPLDLPALIIQRLHGLSAFSLHSPSEINRVASCLRKLGVPLRAWRHRARAVRAKLIQGGRAAICGIYL
FNWAVKTKRKLTPADADRLDLSSWFTVGAGGGDIYHSMSRARPRYLL

GT-6a

MSMSYSWTGALITPCAAEEEEKLPINPLSNLSLRHHNMVYSTTSRSASLRQKKVTFDRVQVFDQHYQDVLKEIKLRA
STVQAKLLSIEEACDLTPSHSARSKYGYGAQDVRSHASKAVNHRSVWEDLLEDSDTPIPTTIMAKNEVFCVDPSK
GGRKPARLIVYPDLGVRVCEKMALYDVTRKLPQAVMGSAAYGFQYSPNQRVEYLLKMWRSKKVPMPGFSYDTRCF

DSTVTERDIRTENDIYQSCQLDPVARRAVSSLTERLYVGGPMVNSKGGQSCGYRRCRASGVLPTSMGNTLTCYLK
AQAACRAANIKDCDMLVCGDDLVICESAGVQEDTASLRAFTDAMTRYSAAPPDAPQPTYDLELITSCSSNVSVA
HDGNGKRYYYLTRDCTTPLARAAWETARHTPVNSWLGNIIMFAPTIWVRMVLMTFFFILQSQEQLKALDFDIYG
VTYSVSPLDLPALIIQRLHGMAAFSLHGYSVELNRVGACLRKLGVPPLRAWRRHRARAVRAKLIQAQGGKAAICGKYL
FNWAVKTKLKLTPVLSASKLDLSGWFVAGYDGGDIYHSVSQARPRLLL