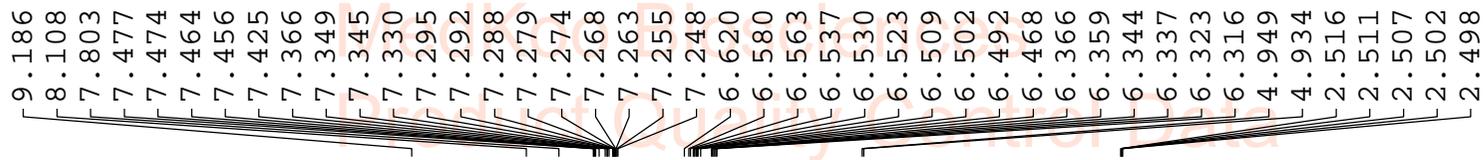


RGFP966 HNMR Analysis, solvent: DMSO-d6



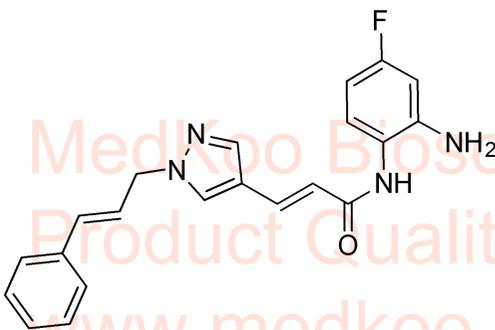
Current Data Parameters
 NAME RGPF-966
 EXPNO 10
 PROCNO 1

F2 - Acquisition Parameters

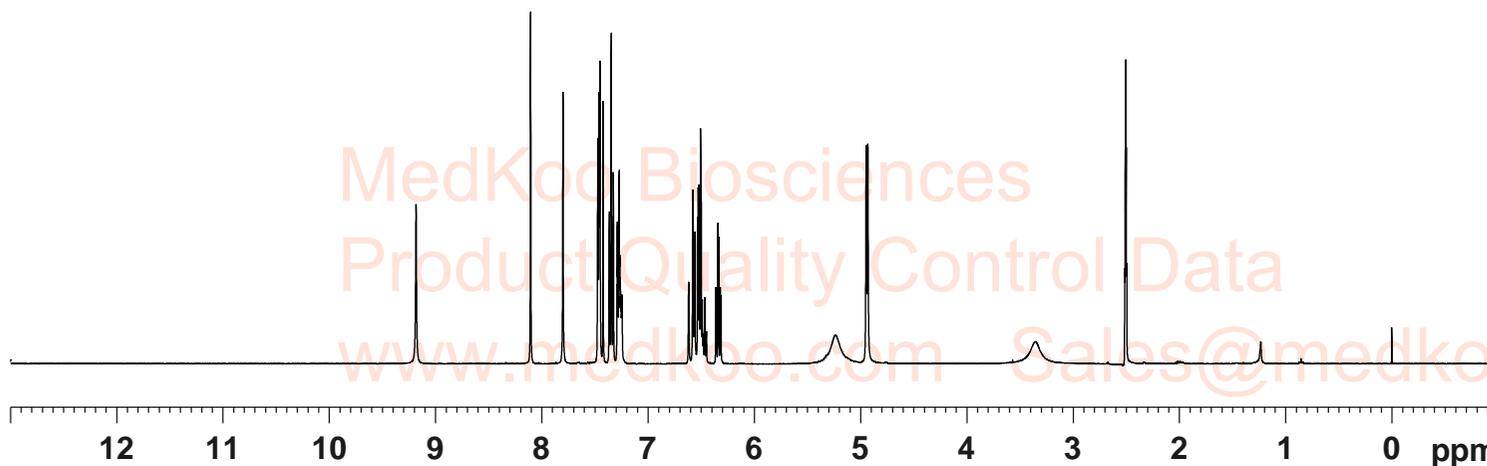
Time 11.10
 INSTRUM spect
 PROBHD 5 mm PABBO BB-
 PULPROG zg30
 TD 65536
 SOLVENT DMSO
 NS 16
 DS 2
 SWH 8012.820 Hz
 FIDRES 0.122266 Hz
 AQ 4.0894465 sec
 RG 161
 DW 62.400 usec
 DE 6.50 usec
 TE 295.7 K
 D1 1.00000000 sec
 TD0 1

==== CHANNEL f1 =====
 SFO1 400.1324710 MHz
 NUC1 1H
 P1 13.00 usec
 PLW1 19.00000000 W

F2 - Processing parameters
 SI 65536
 SF 400.1300000 MHz
 WDW no
 SSB 0
 LB 0 Hz
 GB 0
 PC 1.00

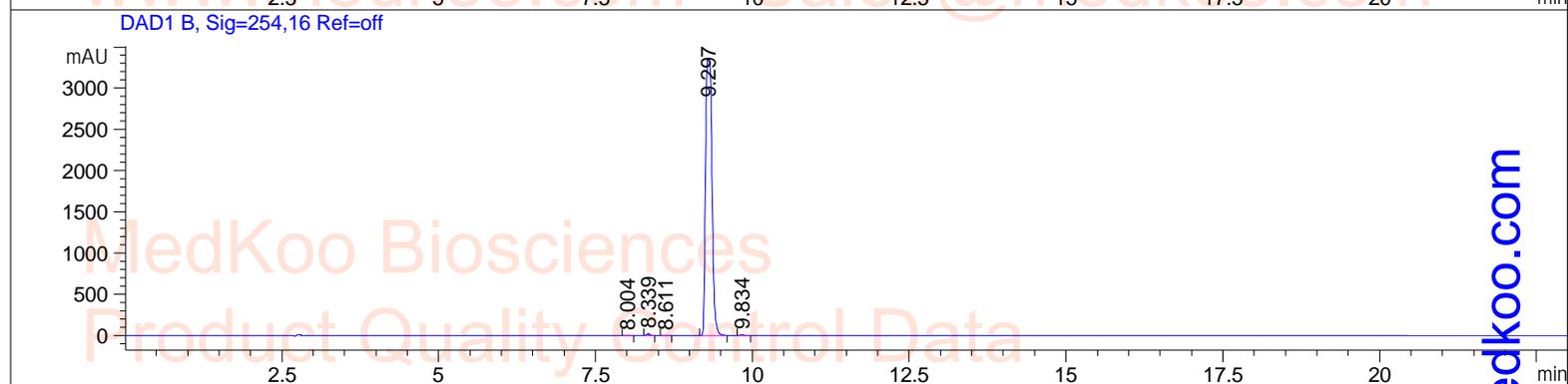
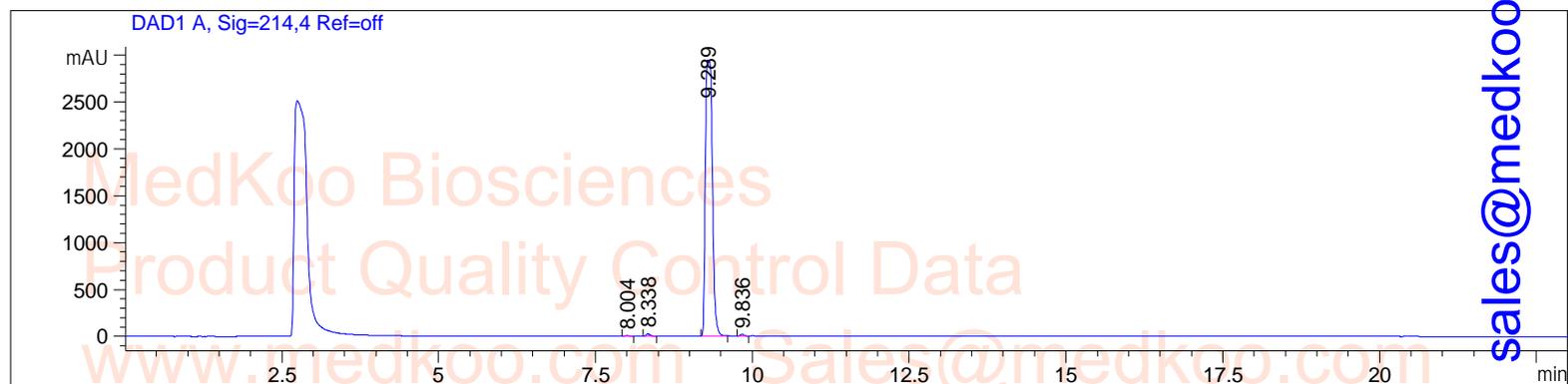


MedKoo Cat.#: 510215
 Name: RGFP966
 CAS#: 1357389-11-7
 Lot#: BBC70328
 Chemical Formula: C₂₁H₁₉FN₄O
 Exact Mass: 362.15429
 Molecular Weight: 362.40



RGFP966 HPLC Analysis

User : Squeunce : 6
 Instrument : Instrument 1 Local : Vial 9
 Method : C:\CHEM32\1\METHODS\TEST-KR.M Injection : 1
 Injection Volume : 2.0 µl



Integration Results

Signal 1: DAD1 A, Sig=214,4 Ref=off

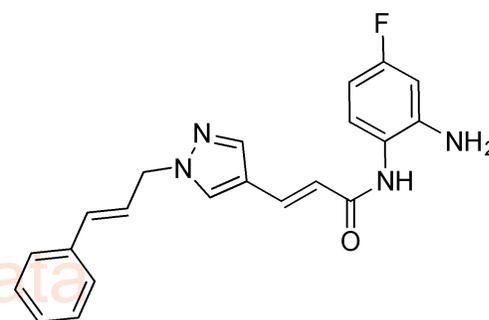
Peak #	Ret. Time [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	8.004	BB	0.0623	31.58629	7.96383	0.1363
2	8.338	BB	0.0667	114.15726	26.29180	0.4927
3	9.289	BB	0.1069	2.29419e4	2943.54712	99.0246
4	9.836	BV	0.0648	80.22608	19.18332	0.3463

Total : 2.31678e4 2996.98607

Signal 2: DAD1 B, Sig=254,16 Ref=off

Peak #	Ret. Time [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	8.004	BB	0.0631	19.21323	4.75579	0.0793
2	8.339	BB	0.0612	87.29830	21.56443	0.3604
3	8.611	BB	0.0601	17.60237	4.65435	0.0727
4	9.297	BB	0.1179	2.40597e4	3346.61792	99.3318
5	9.834	BB	0.0610	37.74636	9.78222	0.1558

Total: 2.42215e4 3387.37470



MedKoo Cat.#: 510215

Name: RGFP966

CAS#: 1357389-11-7

Lot#: BBC70328

Chemical Formula: C₂₁H₁₉FN₄O

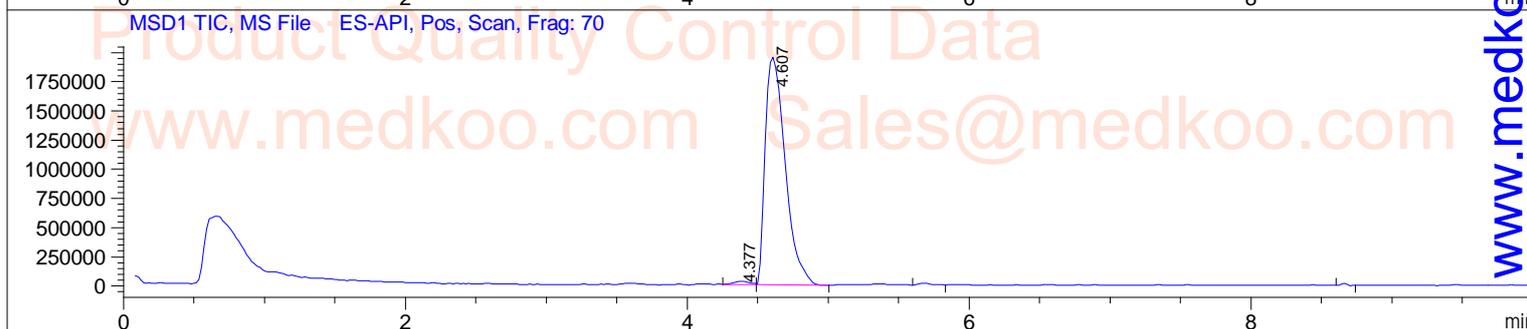
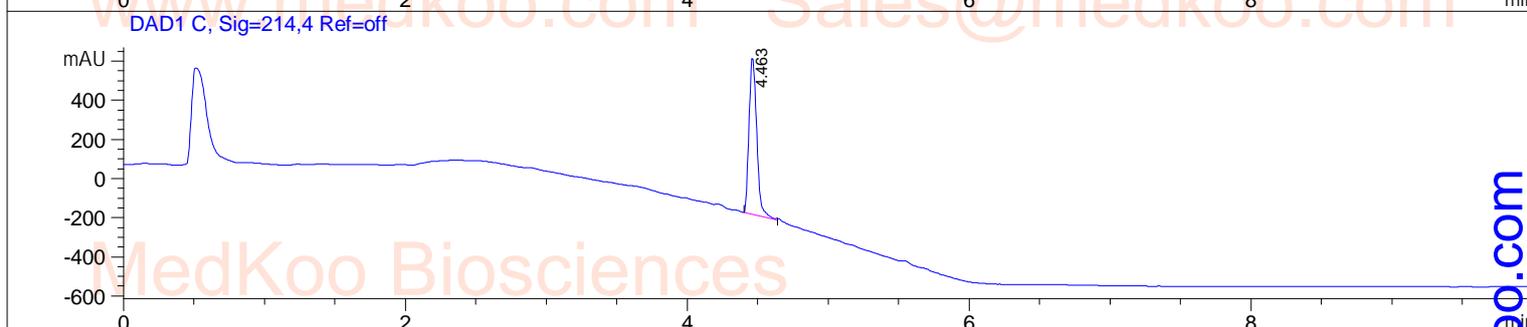
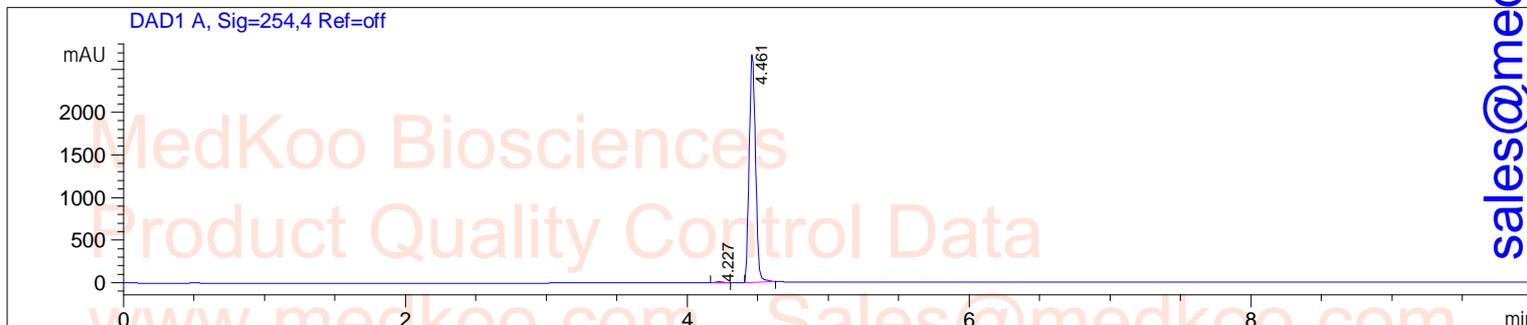
Exact Mass: 362.15429

Molecular Weight: 362.40

RGFP966 LC/MS Analysis

Sample Name : RGPF-966
Acq. Operator : [BSB3]
Spec. Reported : MS Integration
Acq. Method : C:\Chem32\1\DATA\0327\0320 03-27 09-29-27\TEST KR POS.M
Analysis Method : C:\CHEM32\1\METHODS\TEST KR NEG.M

Tgt Mass(EZX) :
Location : Vial 42
Inj : 1
Inj Volume : 1 ul



Ret. Time: 4.61 <<<< POSITIVE SPECTRA >>>>

