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Synthesis, 3D-structure and stability analyses of NRPa-308, a new promising anti-cancer agent.

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ABSTRACT

We report herein the synthesis of a newly described anti-cancer agent, NRPa-308. This compound antagonizes Neuropilin-1, a multi-partners transmembrane receptor overexpressed in numerous tumors, and thereby validated as promising target in oncology. The preparation of NRPa-308 proved challenging because of the orthogonality of the amide and sulphonamide bonds formation. Nevertheless, we succeeded a gram scale synthesis, according to an expeditious three steps route, without intermediate purification. This latter point is of utmost interest in reducing the ecologic impact and production costs in the perspective of further scale-up processes. The purity of NRPa-308 has been attested by means of conventional structural analyses and its crystallisation allowed a structural assessment by X-Ray diffraction. We also reported the remarkable chemical stability of this molecule in acidic, neutral and basic aqueous media. Eventually, we observed for the first time the accumulation of NRPa-308 in two types of human breast cancer cells MDA-MB231 and BT549.

Keywords :

Anti-cancer agents; Neuropilins; NRPa-308 synthesis; NRP-a308 stability

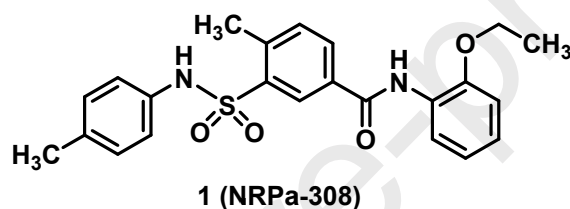
Tumor neo-angiogenesis supplies cancer cells in oxygen and nutrients. Moreover, the neo-formed blood vessels promote also the dissemination of malignant cells to healthy tissues. Therefore, tackling angiogenesis proved to be a relevant therapeutic option in oncology since more than 30 years.⁽¹⁾

Tumor angiogenesis results from the over-expression of specific endothelial cell growth factors, among them the pro-angiogenic isoforms of the Vascular Endothelial Growth Factor (*e.g.* VEGF-A₁₆₅), which bind simultaneously to the tyrosine kinase receptors VEGF-R1 or VEGF-R2, and to neuropilins (NRPs).⁽²⁾ NRPs are multi-partners transmembrane proteins with a non-catalytic cytosolic domain. Although NRPs have been initially described for binding the semaphorins and for their role in neuronal guidance, their involvement in tumor aggressiveness, angiogenesis, lymphangiogenesis and immune escape is now evidenced.⁽³⁾ Moreover, NRPs overexpression is nowadays clinically related to a poor prognosis.

The currently marketed anti-angiogenic drugs, such as Avastin® (a monoclonal antibody directed towards VEGF-A) and Sunitinib® (an ATP mimic targeting the cytosolic domain of tyrosine-kinase involved in angiogenesis), prevents the interaction between VEGF-A₁₆₅ and its receptors VEGF-R1 or VEGF-R2 or directly inhibits their kinase activity, respectively. However, despite indisputable transient benefits for patients, these therapies are not curative; tumors always relapse and become more aggressive, highlighting the real need for alternative therapeutic strategies.⁽⁴⁾ Thereby, in the continuation of our ongoing research in the development of new and potent anticancer agents,⁽⁵⁾ we have focused our attention on NRPs as targets,⁽⁶⁾ and we have recently disclosed NRPa-308 (Chart 1, Compound 1), a small-sized antagonist of the interaction between VEGF-A₁₆₅ and NRPs.⁽⁷⁾ NRPa-308 exerts remarkable anti-angiogenic and anti-proliferative effects *in vitro* (IC₅₀ in the 10 nM range against a large panel of solid and hematological malignancies). Moreover, in our experimental models of nude mice xenografted with human triple negative breast cancer cells (MDA-MB231), NRPa-308

reduces the tumor growth by more than 60%, and enhances significantly animal survival. In addition, this molecule has no acute toxicity in treated animals.⁽⁷⁾

Altogether, these results underline the high potential of NRPa-308 for opening new avenues in anticancer strategies. Thus, we report herein a straightforward synthesis allowing a gram scale production of this molecule. Interestingly, this optimized process does not require intermediate time-consuming purification steps. We also demonstrate the chemical stability of this promising anticancer agent and its significant accumulation in two types of human breast cancer cells (MDA-MB231 and BT549).



NRPa-308 anti-proliferative activity :

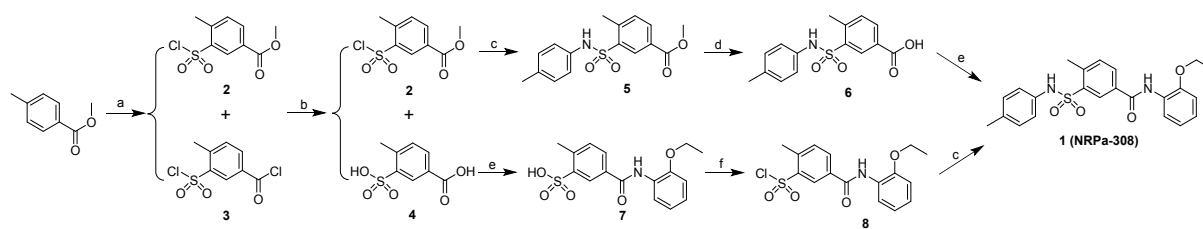
IC₅₀ (48h)	MDA-MB231 : 4.9 μM	IC₅₀ (72h)	MDA-MB231 : 0.2 μM
	BT549 : 2.1 μM		BT549 : 0.1 μM

Sunitinib anti-proliferative activity :

IC₅₀ (48h)	MDA-MB231 : 2.6 μM	IC₅₀ (72h)	MDA-MB231 : 3.7 μM
	BT549 : 2.2 μM		BT549 : 3.5 μM

Chart 1. Structure of NRPa-308, and its antiproliferative activity against two breast cancer cell lines (MDA-MB31 and BT549) measured after 48h and 72h treatment and compared to these of the marketed drug Sunitinib®. These values have been reported by us in ref. 7.

Although NRPa-308 may be described as a “simple” molecule (MW = 424 g/mol, three aromatic rings connected by sulphonamide and amide linkages), it is noteworthy that its large-scale synthesis faces difficulties. This challenge is mainly due to similar synthetic pathways for accessing amide and sulphonamide bonds. Therefore, we decided to smoothly condense successively the required substituted anilines to the central ring, using the commercially available methyl *p*-toluate as starting material.



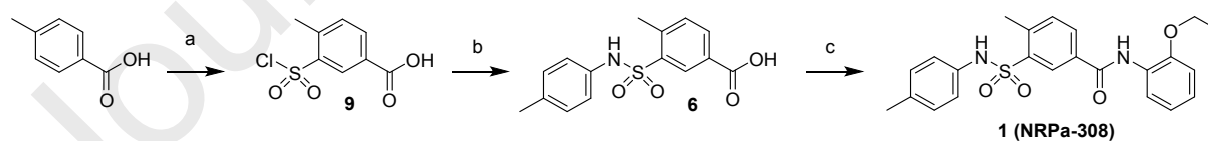
Scheme 1. Reagents and conditions (a): ClSO_3H , 120°C , 4h; (b): 2N aq. HCl , $n\text{Bu}_4\text{N}^+\text{Br}^-$, toluene, reflux overnight; (c): *p*-toluidine, DIEA, DMAP, CH_3CN , R.T. overnight; (d) 2N aq. NaOH , MeOH , 3h, R.T.; (e): 2-ethoxyaniline, BOP, DIEA, DMF , R.T. overnight; (f): SOCl_2 , DMF , R.T., overnight.

Our initial approach is outlined in Scheme 1; the first step consists in the sulfonylation of the methyl *p*-toluate. In fact, despite its apparent chemical simplicity, this reaction is poorly exploited due to its experimental difficulty.⁽⁸⁾ To illustrate this paradox, one can mention the recent work of Singh and co-workers who reported a three-steps process of *p*-toluate sulfonylation through a time-consuming pathway, including successive saponification and re-esterification.⁽⁹⁾ However, we decided to react commercially available methyl *p*-toluate with one equivalent of chlorosulfonic acid under solvent-free conditions to intend the mono-sulfonyl chloride derivative **2**. Different experimental conditions have been tested. We studied the influence of the load of chlorosulfonic acid (from 1.5 eq. until 0.5 eq.) and the influence of the temperature (from 50°C to 130°C , since methyl *p*-toluate melts at 35°C). The best conversion was observed by heating methyl *p*-toluate with 1.1 eq. of chlorosulfonic acid at 120°C for 4 hrs. Nevertheless, this sulfonylation has always led to a mixture of compounds difficult to separate, which consisted of: (i) the expected sulfonyl chloride product **2**; (ii) the dichlorinated compound **3**, and (iii) the remaining unreacted methyl *p*-toluate. With these optimized conditions, the analytic monitoring allowed to estimate the ratio methyl *p*-toluate/**2/3** as about 1/1/3.

This crude material was hydrolysed overnight using 2N aqueous HCl in refluxing toluene with tetrabutylammonium bromide as a phase-transfer catalyst.⁽¹⁰⁾ Although this method is known to hydrolyse selectively a carboxyl chloride *vs* a sulfonyl chloride, this was not the case here since a part of compound **2** has been fully hydrolysed, affording the carboxyl-sulfonic diacid **4**,

which was isolated from the aqueous layer (49% yield). On the other hand, evaporation of the remaining organic layer provided a mixture of a solid containing compound **2** and the unreacted methyl *p*-toluate (20% yield), and compound **3** as an oil. The two layers were separated by filtration. Unpurified compound **2** was coupled with *p*-toluidine using catalytic amounts of DMAP to afford sulfonamide **5**,⁽¹¹⁾ which was then saponified to provide **6** after acidification. The *p*-toluic acid resulting from the unreacted methyl *p*-toluate was removed at this step by repeated trituration in diethyl ether affording compound **6** as a pure white solid (31% yield). The coupling of **6** with 2-ethoxyaniline using BOP/DIEA in DMF led to the final expected compound **1** (NRPa-308) in 73% yield.

According to this process, the recycling of **4** provided an alternative route to compound **1** (Scheme 1). Indeed, compound **4** was treated with *o*-ethoxyaniline in presence of BOP/DIEA. The resulting compound **7** was obtained in 62% yield, and treated with thionyl chloride at room temperature;⁽¹²⁾ it afforded a chloro-sulfonic acid derivative, which was directly condensed with *p*-toluidine in the presence of catalytic DMAP. Following this pathway, compound **1** was obtained in 24% yield (over two steps). Although this process does not require any purification step, which is very important to speed up a synthesis, it does not allow any scale-up, since the desired product was obtained in less than 10% yield from the commercial starting material.



Scheme 2. Reagents and conditions (a): ClSO_3H , 100°C , 12h; (b): *p*-toluidine, DCM, 0°C - R.T. 1h; (c): *o*-ethoxyaniline, EDCI, DIEA, DMF, R.T. overnight

Therefore, we have considered the alternative synthetic route outlined in scheme 2. According to this second pathway, the target compound **1** may be obtained in only three steps from *p*-methylbenzoic acid as commercially available starting material. The chlorosulfonylation has

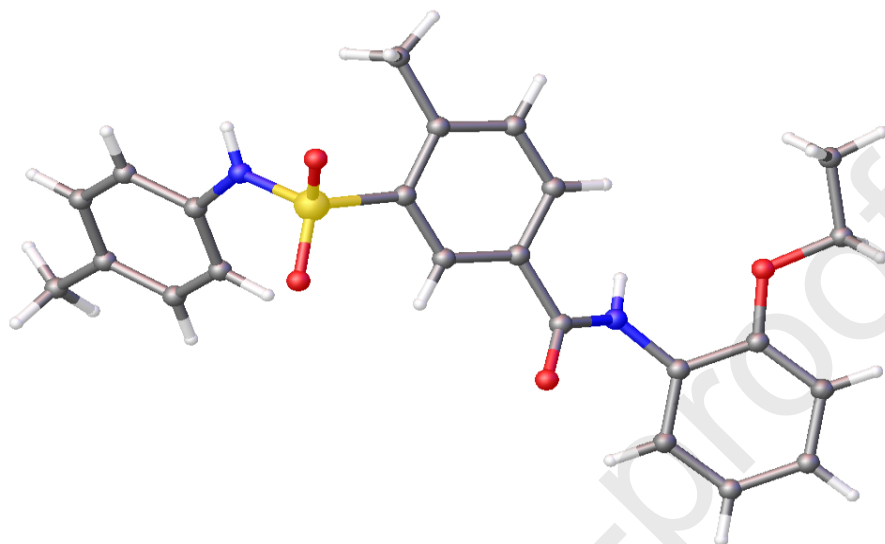
been conveniently performed by heating *p*-methylbenzoic acid in chlorosulfonic acid at 100°C. The reaction occurs overnight, however the subsequent work-up requires a sustained attention. Indeed, the mixture should be cautiously ice-quenched to avoid an exothermic behaviour leading to the formation of side-products. After the neutralization of the unreacted chlorosulfonic acid, conventional filtration and water washing afforded **9** as a white solid in 84% yield. The unpurified product was directly reacted at 0°C in dichloromethane with *p*-toluidine for one hour to afford the bi-aryl sulfonamide **6**. The expected product was isolated as a pure solid after acidic treatment and extraction (78% yield). ¹H NMR provided evidence that no purification was required at this step. Eventually, compound **6** has been reacted according to a conventional process with *o*-ethoxyaniline (1.1 eq.) in the presence of EDCI (1.1 eq) as coupling agent and diisopropylethylamine (3.1 eq.) as base. This last reaction has been completed overnight, and it afforded after treatment and purification by flash chromatography the expected NRPa-308 (**1**) as a pure white solid (64% yield).

To summarize, this second synthetic route afforded NRPa-308 **1** in shortened reaction times and at the gram scale (global yield from the commercially available starting material: 42%). Another key feature is the absence of intermediate purifications, which reduces the ecologic impact and the production costs of the synthesis (no use of large amount of toxic solvents requiring recycling).

In addition, single crystals of NRPa-308 **1** were obtained from hot toluene, and a suitable one was selected for X-Ray 3-D structure determination. The experimental procedure is depicted in the Supporting Information section. Briefly, the structure was solved by direct method using SHELXS⁽¹³⁾ refinement, based on F² was carried out by full matrix least squares using SHELXL-2018⁽¹⁴⁾ software with anisotropic displacement parameters for all non-hydrogen atoms. Hydrogen atoms were located on a difference Fourier map and introduced into that

calculations as a riding model with isotropic thermal parameters. All calculations were performed by using the Crystal Structure crystallographic software package WINGX.⁽¹⁵⁾

A



B

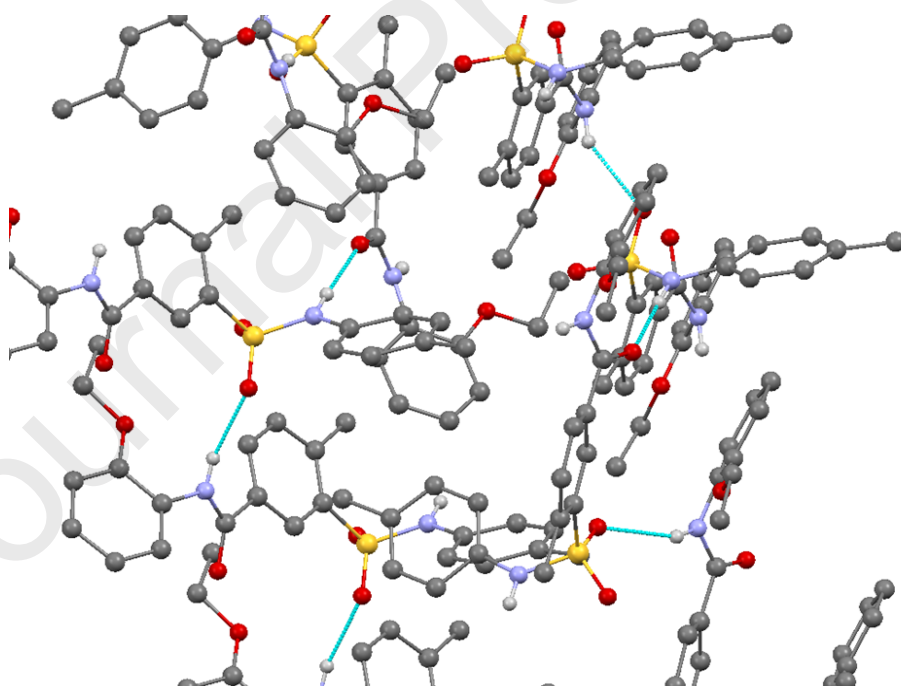


Figure 1. Structure of NRPa-308 **1** analysed by X Ray. Please refer to reference 16, and to the Supporting Information section for more details. **A:** Single crystal of Nrp-a308 **1**; **B:** Supramolecular arrangement of Nrp-a308 **1**.

NRPa-308 **1** has a molecular structure built from entities, depicted in Figure 1.⁽¹⁶⁾ The crystal data collection and refinement parameters are collected in Table S1 (Supporting Information

section). CCDC-1939102 contains the supplementary crystallographic data for this paper, which can be obtained free of charge from the Cambridge Crystallographic Data centre.

Briefly, at the single molecular level, the two aromatic rings linked thanks to the amide bond appear rather coplanar, while the third one, connected by the sulphonamide linker, is twisted, rather perpendicular to the central aromatic ring. At the supramolecular architecture level, neutral molecules are associated in the crystal essentially via N-H \cdots O hydrogen bonds in a three-dimensional way. More precisely, two hydrogen bonds involving three close molecules are evidenced in this structure: (i) the first involves the NH “amide” of the “central” molecule, and an oxygen of the sulfoxide linker belonging to an adjacent molecule; (ii) the second takes place between the NH “sulfonamide” of the “central” molecule and the oxygen of the amide belonging to a third molecule. In the continuation of this structural analysis, we plan now to crystalize NRPa-308 **1** with NRP-1 to decipher the close contacts between the antagonist and its receptor.

We next focused our attention on NRPa-308 **1** chemical stability. Interestingly, we reported formerly that the anti-proliferative activities of NRPa-308 are deeply related to its incubation times with the tumor cells (Chart 1).⁽⁷⁾ Thus, our purpose was to unambiguously demonstrate that these anti-proliferative effects are solely due to NRPa-308, and not to one of its potential metabolites or one of its degradation products.

First, the chemical stability of NRPa-308 was assayed by HPLC analysis at three different pH (0.9, 7.4 and 8.4) and at two temperatures (25°C and 37°C). After 16 days, the compound proved fully stable under all these conditions (**Figure 2**, the detailed experimental procedures are given in the Supporting Information section).

Next, we studied the stability of NRPa-308 in cellular culture media, in presence of malignant cells. To this end, MDA-MB-231 and BT-549 breast cancer cells were cultured in the presence

of 2 μ mol of NRPa-308 for three days, and the HPLC analyses of the supernatant revealed that NRP-a308 was not degraded (Figure S1, Supporting Information section). Then, the cells were washed and lysed with methanol. The lysates were extracted with $\text{CHCl}_3/\text{MeOH}$, 9/1, v/v and quantitatively analysed by HPLC. In both cases, NRPa-308 proved stable, with no other peak detected (**Figure 3**, the detailed experimental procedures are given in the Supporting Information section). Therefore, the *in vitro* antiproliferative activities, already measured by us (Chart 1), are due to compound **1** and not due to one of its metabolites or degradation products.

In addition, the quantitative dosage revealed a significant accumulation of NRPa-308 in cells, with respectively $5.14 \pm 0.06 \cdot 10^{-9}$ mol and $1.40 \pm 0.01 \cdot 10^{-9}$ mol calculated in MDA-MB231 and BT549 cells, after three days. Indeed, NRP-a308 mimics the interaction between NRP-1 and the endogenous tetrapeptide TKPR, so-called tuftsin. Tuftsin has been used as model for the development of penetrating peptides able to interact with the VEGF- A_{165} /NRP-1 hotspot.^(17b) The tuftsin-derivative penetrating peptides respect the C-terminus sequence R/K/XXR/K (so called : C-end rule or CendR). They interact thereby with NRP-1, and are internalized into the cells thanks to the NRP-1 transmembrane trafficking. Based on this mechanism, peptide carriers (iRGD peptides) for selective drugs delivery have been recently disclosed. The iRGDs carriers are short-sized cyclic peptides, whose cleavage by integrins deliver a “C-end rule” sequence, able to be internalized into cells by NRP-1.⁽¹⁸⁾ The iRGDs are used to selectively address inside malignant cells potential therapeutic agents, such as small-sized molecules (e.g. doxorubicin) or siRNAs.⁽¹⁹⁾ However, at the best of our knowledge, non-peptidic small-sized NRP-1 antagonists have ever been reported for a potential “iRGD” like vectorization process. Thus, this result might constitute a way for the development of a new class of non-peptidic compounds able to selectively address therapeutic agents into the tumor cells through NRP-1.

To conclude, we report herein an expeditious synthesis for the newly identified anti-cancer agent NRPa-308. Our strategy is based on a three steps procedure, and it requires neither intermediate purification nor use of expensive reactants, which is of utmost interest for the development of further large-scale production. The structure of this molecule has been unambiguously characterised thanks to X-Ray crystallography, and the three-dimensional supramolecular architecture of the crystal is granted by N-H...O hydrogen bonds between three adjacent molecules. We also observed the acute stability of this molecule in different media, which proved that its remarkable cytotoxic effects against cancer cells is not imputable to any of its potential metabolites. Lastly, we report the significant accumulation of NRPa-308 into cancer cells, which might be related to the transmembrane trafficking properties of NRP-1.

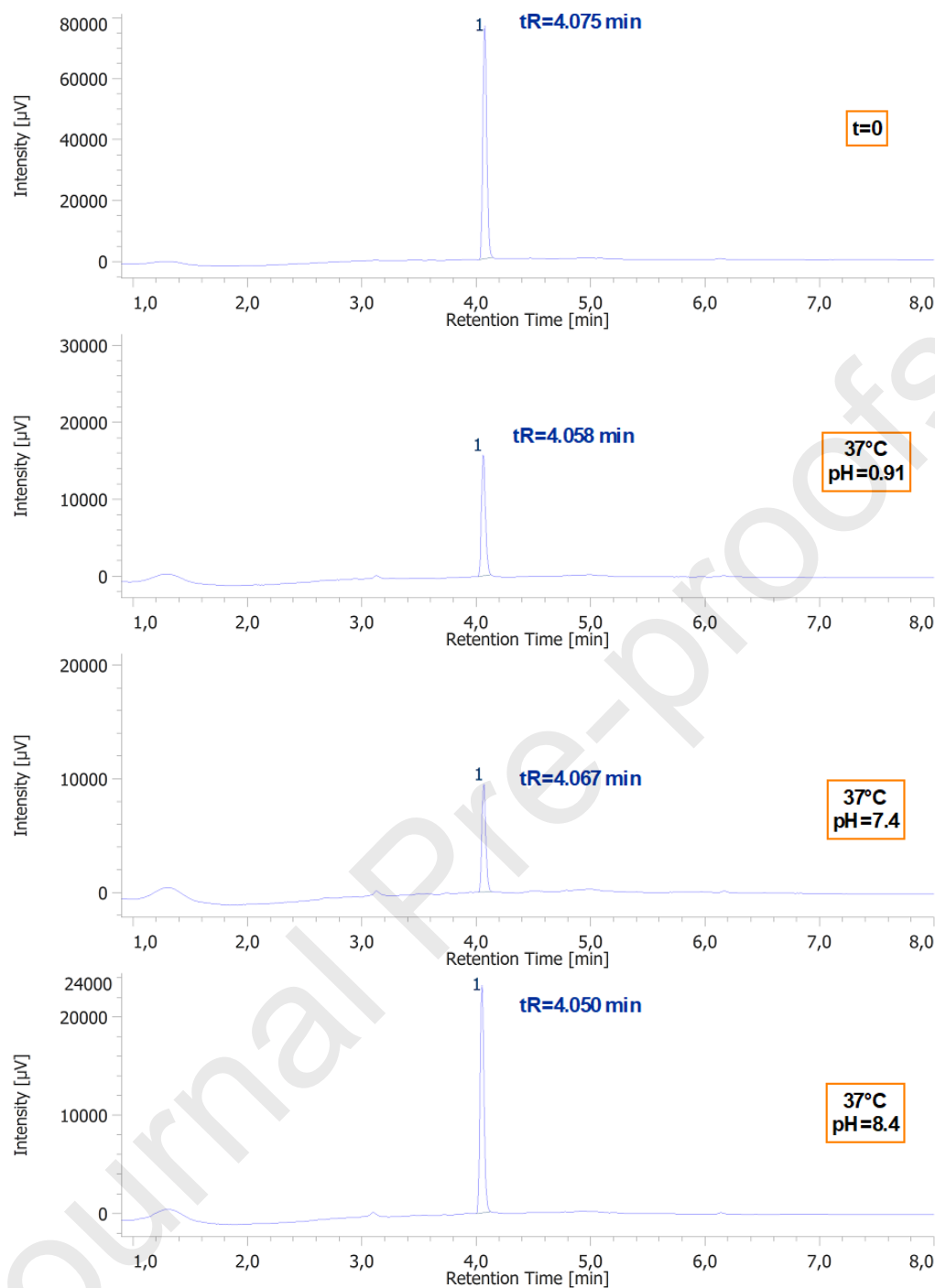


Figure 2. Chromatograms showing the stability of NRPa-308 in aqueous buffers at pH 0.9, 7.4 and 8.4 after 16 days incubation at 37°C.

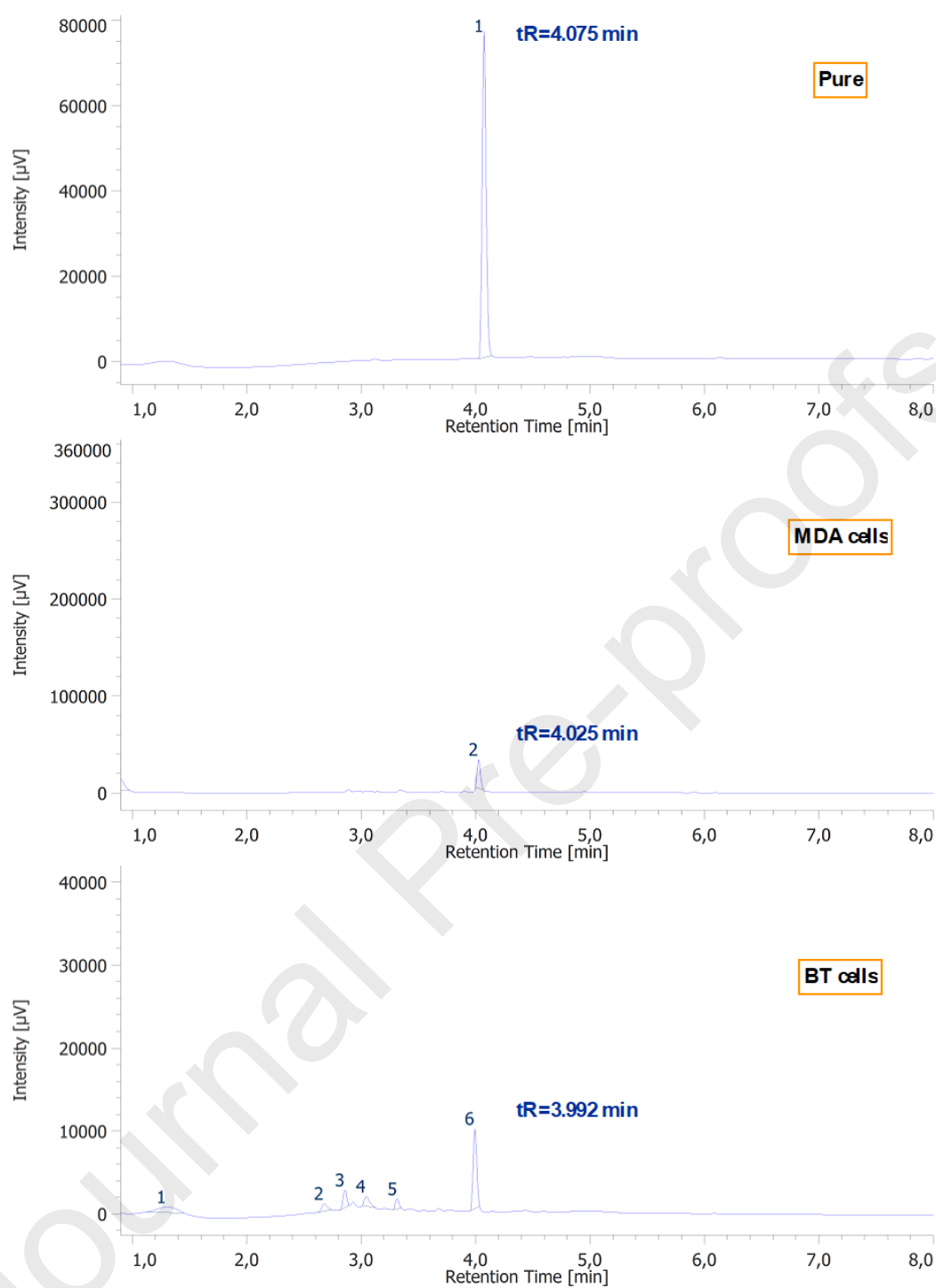
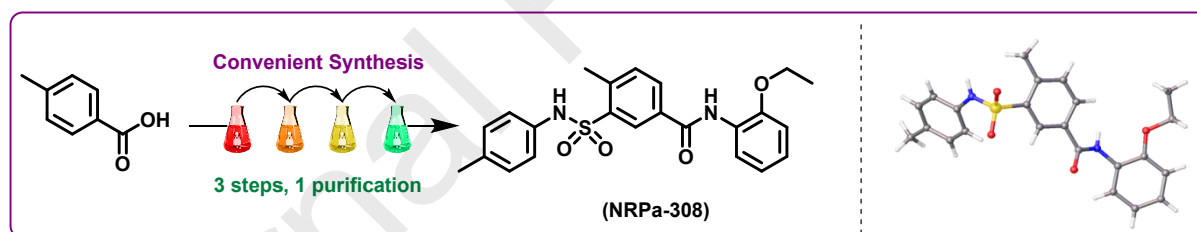


Figure 3. Chromatograms showing the metabolism of NRPa-308 in MDA-MB231 and BT549 cells after three days incubation at 37°C.

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Graphical abstract



References

1. a) Arezumand R, Alibakhshi A, RanjbariJ, Ramazani A, Muyldermans S. *Frontiers in Immunology* 2017, 8, 1746/1-1746/13; b) Martin AR, Ronco C, Demange L, Benhida R. *Med. Chem. Commun.* 2017, 8, 21-52; c) Frandsen S, Kopp S, Wehland M, Pietsch J, Infanger M, Grimm D. *Curr. Pharm. Design* 2016, 22, 5927-5942.

2. a) Bergantino F, Guariniello S, Raucci R, Colonna G, DeLuca A, Normanno N, Costantini S. *Biochim. Biophys. Acta* 2015, 1854, 410-425; b) Horowitz A, Seerapu HR. *Cell. Signal.* 2012, 24, 1810-1820 ; c) Ferrara N, Gerber HP, LeCouter J. *Nature Med.* 2003, 9, 669-676.
3. a) Roy S, Bag AK, Singh RK, Talmadge JE, Batra SK, Datta K. *Front. Immunol.* 2017, 8, 1228/1-1228/27; b) Guo HF, Vander Kooi CW *J. Biol. Chem.* 2015, 290, 29120-2916; b) Prud'homme GJ, Glinka Y. *Oncotarget* 2012, 3, 921-939; c) Jubb AM, Strickland L, Liu SD, Mak J, Schmidt M, Koeppen H. *J. Pathol.* 2012, 226, 50-60; d) Koch S. *Biochem. Soc. Trans.* 2012, 40, 20-25; e) Grandclement C, Borg C. *Cancers* 2011, 3, 1899-1928.
4. a) Spanheimer PM, Lorenzen AW, De Andrade JP, Kulak MV, Carr JC, Woodfield GW, Sugg SL, Weigel RJ. *Ann. Surg. Oncol.* 2015, 22, 4287-4294; b) Gaumann AK, Kiefer F, Alfer J, Lang SA, Geisser EK, Breier G. *Int. J. Cancer* 2016, 138, 540-554; c) Giuliano S, Cormerais Y, Dufies M, Grepin R, Colosette P, Belaid A, Parola J, Martin A, Lacas Gervais S, Mazure NM, Benhida R, Auberger P, Mograbi B, Pages G. *Autophagy* 2015, 11, 1891-1904.
5. a) Dufies M, Grytsai O, Ronco C, Camara O, Ambrosetti D, Parola J, Lagarde N, Montès M, Mateo L, Giuliano S, Grépin R, Auberger P, Demange L, Pagès G, Benhida R. *Theranostics* 2019, doi: 10.7150/thno.35032 ; b) Marzag H, Zerhouni M, Tachallait H, Demange L, Robert G, Bougrin K, Auberger P, Benhida R. *Bioorg. Med. Chem. Lett.* 2018, 28, 1931-1936; c) Allaoui S, Dufies M, Driowya M, Demange L, Bougrin K, Robert G, Auberger P, Pagès G, Benhida R. *Bioorg Med. Chem. Lett.* 2017, 27(9), 1989-1992; d) Ronco C, Martin AR, Demange L, Benhida R. *Med. Chem. Commun.* 2017, 8(2), 295-319; e) Houzé S, Hoang NT, Lozach O, LeBras J, Meijer L, Galons H, Demange L. *Molecules* 2014, 19(9), 15237-15257.
6. a) Jarray R, Pavoni S, Borriello L, Allain B, Lopez N, Bianco S, Liu WQ, Biard D, Demange L, Hermine O, Garbay C, Raynaud F, Lepelletier Y. *Biochimie* 2015, 118, 151-161; b) Liu WQ, Borriello L, Allain B, Pavoni S, Hermine O, Garbay C, Raynaud F, Lepelletier Y, Demange L. *Int. J. Pept. Res. Ther.* 2015, 21(1), 117-124; c) Goldwasser E, DeCourcy B, Demange L, Garbay

- C, Raynaud F, Hadj-Slimane R, Piquemal JP, Gresh N. *J. Mol. Model.* 2014, 20(11), 1-24; d) Borriello L, Montès M, Lepelletier Y, Leforban B, Liu WQ, Demange L, Delhomme B, Pavoni S, Jarray R, Boucher JL, Dufour S, Hermine O, Garbay C, Hadj-Slimane R, Raynaud F. *Cancer Lett.* 2014, 349(2), 120-127; e) Liu WQ, Megale V, Borriello L, Leforban B, Montès M, Goldwasser E, Gresh N, Piquemal JP, Hadj-Slimane R, Hermine O, Garbay C, Raynaud F, Lepelletier Y, Demange L. *Bioorg. Med. Chem. Lett.* 2014, 24(17), 4254-4259.
7. Liu WQ, Lepelletier Y, Montes M, Borriello L, Jarray R, Grépin R, Leforban B, Loukaci A, Benhida R, Hermine O, Dufour S, Pagès G, Garbay C, Raynaud F, Hadj-Slimane R, Demange L. *Cancer Lett.* 2018, 414, 88-98.
8. a) Abbavaram BRA, Reddyvari HRV. *J. Kor. Chem. Soc.* 2013, 57, 731-737; b) Kendall JD, Giddens AC, Tsang KY, Frédérick R, Marshall ES, Singh R, Lill CL, Lee WJ, Kolekar S, Chao M, Malik A, Yu S, Chaussade C, Buchanan C, Rewcastle GW, Baguley BC, Flanagan JU, Jamieson SMF, Denny WA, Shepherd PR. *Bioorg. Med. Chem.* 2012, 20, 58-68; c) Zhang R, Lei L, Xu YG, Hua WY, Gong GQ. *Bioorg. Med. Chem.* 2007, 17, 2430-2433.
9. Singh R, Bordeaux M, Fasan R. *ACS Catal.* 2014, 4, 546-552.
10. Ashikawa M, Sakagami S. JP 2009167142, 2009.
11. Hadj-Slimane R. WO 2015004212, 2015.
12. Jenkis TJ, Guan B, Dai M, Li G, Lightburn TE, Huang S, Freeze BS, Burdi DF, Jacutin-Porte S, Bennett R, Chen W, Minor C, Ghosh S, Blackburn C, Gigstad KM, Jones M, Kolbeck R, Yin W, Smith S, Cardillo D, Ocain TD, Harriman GC. *J. Med. Chem.* 2007, 50, 566-584.
13. Sheldrick GM. SHELXS-97, Program for Crystal Structure Solution, University of Göttingen, Göttingen, Germany, 1997.
14. Sheldrick GM *Acta Crystallogr., sect. A: Found Crystallogr.* 2008, 64, 112-122.
15. Farrugia LJ. *J. Appl. Cryst.* 1999, 32, 837.

16. Crystal data and structure refinement for NRPa-308: Orthorhombic crystal, space group $P2_12_12_1$ (no. 19), $a = 7.8694(2) \text{ \AA}$, $b = 14.9423(5) \text{ \AA}$, $c = 17.9015(5) \text{ \AA}$, $V = 2104.98(11) \text{ \AA}^3$, $Z = 4$, $T = 100(2) \text{ K}$, $\mu(\text{CuK}\alpha) = 1.597 \text{ mm}^{-1}$, $D_{\text{calc}} = 1.339 \text{ g/cm}^3$, 23575 reflections measured ($7.706^\circ \leq 2\theta \leq 133.216^\circ$), 3719 unique ($R_{\text{int}} = 0.0396$, $R_{\text{sigma}} = 0.0264$) which were used in all calculations. The final R_1 was 0.0252 ($I > 2\sigma(I)$) and wR_2 was 0.0617 (all data). More details are provided in the Supporting Information section.

17. a) Zanut D, Kotla R, Nussinov R, Teesalu T, Sugahara KN, Aleman C, Haspel N. *J. Struct. Biol.* 2013, 182, 78-86 b) Teesalu T, Sugahara KN, Kotamraju VR, Ruoslahti E. *Proc. Natl. Acad. Sci. USA* 2009, 106, 16157-16162.

18. a) Roth L, Agemy L, Kotamraju VR, Braun G, Teesalu T, Sugahara KN, Hamzah J, Ruoslahti E. *Oncogene* 2012, 31, 3754-3763; b) Ye Y, Zhu L, Ma Y, Niu G, Chen X. *Bioorg Med. Chem. Lett.* 2011, 21, 1146-1150; c) Fu KF, Zhang WQ, Luo LM, Song P, Li D, Du R, Ren W, Huang D, Lu WL, Zhang X, Zhang Q. *Int. J. Nanomed.* 2013, 8, 2473-2485; d) Sugahara KN, Teesalu T, Karmali PP, Kotamraju VR, Agemy L, Girard OM, Hanahan D, Mattrey RF, Ruoslahti E. *Cancer Cell* 2009, 16, 510-520.

19. a) Liu H, Shi X, Wu D, Khshen FK, Deng L, Dong A, Wang W, Zhang J *ACS Appl. Mater. Interfaces* 2019, 11, 19700-19711; b) Ding N, Zou Z, Sha H, Su S, Qian H, Meng F, Chen F, Du S, Zhou S, Chen H, Zhang L, Yang J, Wei J, Liu B *Nature Commun.* 2019, doi: 10.1038/s41467-019-09296-6; c) Barman S, das G, Gupta V, Mondal P, Jana B, Bhunia D, Khan J, Mukherjee D, Ghosh S *Mol. Pharmaceutics* 2019, 16, 2522-2531; d) Lo JH, Hao L, Muzumdar MD, Raghavan S, Kwon EJ, Pulver EM, Hsu F, Aguirre AJ, Wolpin BM, Fuchs CS, Hahn WC, Jacks T, Bhatia SN *Mol. Cancer Ther.* 2018, 17(11), 2377-2388.

