(12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

CORRECTED VERSION

(19) World Intellectual Property Organization
International Bureau

(43) International Publication Date
6 November 2008 (06.11.2008)

PCT

(10) International Publication Number
WO 2008/132134 A9

(51) International Patent Classification:
C07K 16/14 (2006.01) *C12N 15/10* (2006.01)
A61K 39/395 (2006.01) *C07K 16/18* (2006.01)
A61P 31/00 (2006.01) *A61P 37/06* (2006.01)
A61K 35/00 (2006.01)

(21) International Application Number:
PCT/EP2008/055006

(22) International Filing Date: 24 April 2008 (24.04.2008)

(25) Filing Language: English

(26) Publication Language: English

(30) Priority Data:
07107140.1 27 April 2007 (27.04.2007) EP
07113353.2 27 July 2007 (27.07.2007) EP

(71) Applicant *(for all designated States except US)*: **NOVARTIS AG** [CH/CH]; Lichtstrasse 35, CH-4056 Basel (CH).

(72) Inventors; and

(75) Inventors/Applicants *(for US only)*: **BURNIE, James** [GB/GB]; 1 Gresytoke Drive, Alderley Edge, Cheshire SK9 7PY (GB). **WECHNER, Philipp** [AT/AT]; Hussistrasse 27c, A-6130 Schwaz (AT).

(74) Agent: **GRUBER, Markus**; Novartis Pharma AG, Patent Department, CH-4002 Basel (CH).

(81) Designated States *(unless otherwise indicated, for every kind of national protection available)*: AE, AG, AL, AM, AO, AT, AU, AZ, BA, BB, BG, BH, BR, BW, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DO, DZ, EC, EE, EG, ES, FI, GB, GD, GE, GH, GM, GT, HN, HR, HU, ID, IL, IN, IS, JP, KE, KG, KM, KN, KP, KR, KZ, LA, LC, LK, LR, LS, LT, LU, LY, MA, MD, ME, MG, MK, MN, MW, MX, MY, MZ, NA, NG, NI, NO, NZ, OM, PG, PH, PL, PT, RO, RS, RU, SC, SD, SE, SG, SK, SL, SM, SV, SY, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VC, VN, ZA, ZM, ZW.

(84) Designated States *(unless otherwise indicated, for every kind of regional protection available)*: ARIPO (BW, GH, GM, KE, LS, MW, MZ, NA, SD, SL, SZ, TZ, UG, ZM,

[Continued on next page]

(54) Title: NOVEL ANTIBODY MOLECULES AND NUCLEIC ACIDS BINDING TO FUNGAL STRESS PROTEIN HSP90

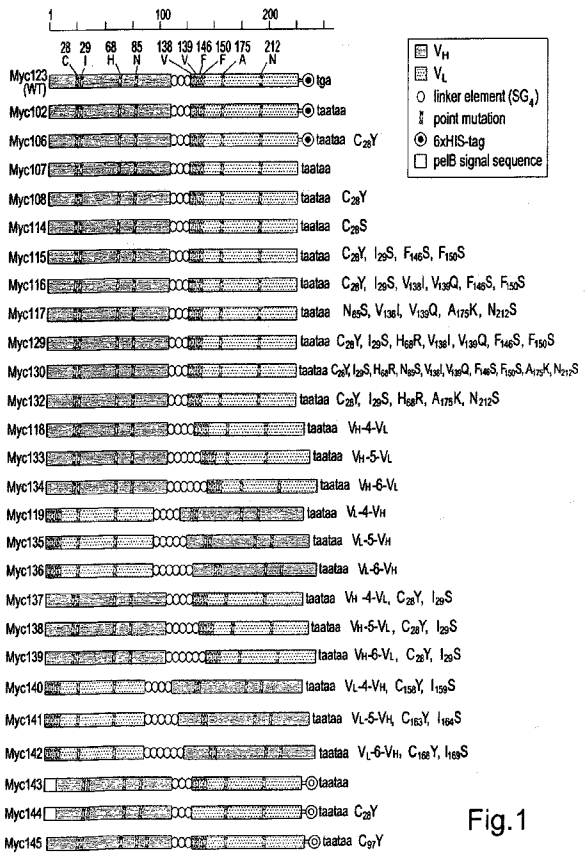


Fig.1

(57) Abstract: An scFv peptide comprising a V_H domain and a V_L domain linked by an amino acid spacer is disclosed. The V_H domain comprises a sequence with at least 80% sequence identity to the sequence of SEQ ID NO. 64. The V_L domain comprises a sequence with at least 80% sequence identity to the sequence of SEQ ID NO. 66. The scFv peptide also comprises the substitution or deletion of an amino acid in the V_H domain at the position corresponding to C_{28}.

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ZW), Eurasian (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European (AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HR, HU, IE, IS, IT, LT, LU, LV, MC, MT, NL, NO, PL, PT, RO, SE, SI, SK, TR), OAPI (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG).

Published:

— *with international search report*

— *with sequence listing part of description published separately in electronic form and available upon request from the International Bureau*

(48) Date of publication of this corrected version:
9 July 2009

(15) Information about Correction:
see Notice of 9 July 2009