

(19) World Intellectual Property
Organization
International Bureau

(43) International Publication Date
01 November 2018 (01.11.2018)



(10) International Publication Number
WO 2018/200380 A8

(51) International Patent Classification:

C12Q 1/6806 (2018.01) C12Q 1/6855 (2018.01)

(21) International Application Number:

PCT/US2018/028867

(22) International Filing Date:

23 April 2018 (23.04.2018)

(25) Filing Language:

English

(26) Publication Language:

English

(30) Priority Data:

62/488,830 23 April 2017 (23.04.2017) US

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(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, BN, BR, BW, BY, BZ, CA, CH, CL, CN, CO, CR, CU, CZ, DE, DJ, DK, DM, DO, DZ, EC, EE, EG, ES, FI, GB, GD, GE, GH, GM, GT, HN, HR, HU, ID, IL, IN, IR, IS, JO, JP, KE, KG, KH, KN, KP, KR, KW, KZ, LA, LC, LK, LR, LS, LU, LY, MA, MD, ME, MG, MK, MN, MW, MX, MY, MZ, NA, NG, NI, NO, NZ, OM, PA, PE, PG, PH, PL, PT, QA, RO, RS, RU, RW, SA, SC, SD, SE, SG, SK, SL, SM, ST, SV, SY, TH, 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, LR, LS, MW, MZ, NA, RW, SD, SL, ST, SZ, TZ, UG, ZM, ZW), Eurasian (AM, AZ, BY, KG, KZ, RU, TJ,

(54) Title: COMPOSITIONS AND METHODS FOR IMPROVING SAMPLE IDENTIFICATION IN INDEXED NUCLEIC ACID LIBRARIES

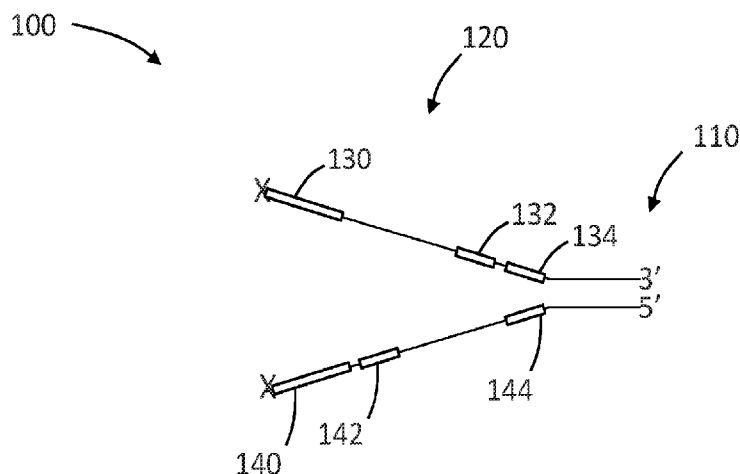


FIG. 1

(57) Abstract: The present invention is concerned with compositions and methods for improving the rate of correct sample identification in indexed nucleic acid library preparations for multiplex next generation sequencing by exonuclease treatment after protective adapters are ligated to target polynucleotides to degrade unincorporated adapters prior to amplification and sequencing.



TM), European (AL, AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HR, HU, IE, IS, IT, LT, LU, LV, MC, MK, MT, NL, NO, PL, PT, RO, RS, SE, SI, SK, SM, TR), OAPI (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, KM, ML, MR, NE, SN, TD, TG).

Declarations under Rule 4.17:

- *as to applicant's entitlement to apply for and be granted a patent (Rule 4.17(ii))*

Published:

- *with international search report (Art. 21(3))*

(48) Date of publication of this corrected version:

05 December 2019 (05.12.2019)

(15) Information about Correction:

see Notice of 05 December 2019 (05.12.2019)