

# Appendix C

## Published works

### Named authorships

Wedge, D. C.\*, Gundem, G.\*, Mitchell, T.\*, Martincorena, I., Ghori, M., Zamora, J., Butler, A., Whitaker, H., Kote-Jarai, Z., Alexandrov, L. B., Van Loo, P., Massie, C. E., **Dentro, S.**, Warren, A. Y., Verrill, C., Berney, D., Dennis, N., Merson, S., Hawkins, S., Howat, W., Yu, Y., Lambert, A., Kay, J., Kremeyer, B., Karaszi, K., Luxton, H., Camacho, N., Marsden, L., Edwards, S., Matthews, L., Bo, V., Leongamornlert, D., McLaren, S., Ng, A., Yu, Y., Zhang, H., Dadaev, T., Thomas, S., Easton, D. F., Ahmed, M., Bancroft, E., Fisher, C., Livni, N., Nicol, D., Tavaré, S., Gill, P., Greenman, C., Khoo, V., Van As, N., Kumar, P., Ogden, C., Cahill, D., Thompson, A., Mayer, E., Rowe, E., Dudderidge, T., Gnanapragasam, V., Shah, N. C., Raine, K., Jones, D., Menzies, A., Stebbings, L., Teague, J., Hazell, S., de Bono, J., Attard, G., Isaacs, W., Visakorpi, T., Fraser, M., Boutros, P. C., Bristow, R. G., Workman, P., Sander, C., The TCGA consortium, Hamdy, F. C., Futreal, A., McDermott, U., Al-Lazikani, B., Lynch, A. G., Bova, G. S., Foster, C. S., Brewer, D., Neal, D., Cooper, C. S., and Eeles, R. A. Sequencing of prostate cancers identifies new cancer genes, routes of progression and drug targets. *Nature Genetics* - Manuscript accepted.

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Gerstung, M.\*, Jolly, C.\*, Leshchiner, I.\*, **Dentro, S. C.\***, Gonzalez, S., Mitchell, T. J., Rubanova, Y., Anur, P., Rosebrock, D., Yu, K., Tarabichi, M., Deshwar, A., Wintersinger, J., Kleinheinz, K., Vazquez-Garcia, I., Haase, K., Sengupta, S., Macintyre, G., Malikic, S., Donmez, N., Livitz, D. G., Cmero, M., Demeulemeester, J., Schumacher, S., Fan, Y., Yao, X., Lee, J., Schlesner, M., Boutros, P. C., Bowtell, D. D., Zhu, H., Getz, G., Imielinski, M., Beroukhim, R., Sahinalp, S. C., Ji, Y., Peifer, M., Markowitz, F., Mustonen, V., Yuan, K., Wang, W., Morris, Q. D., Spellman, P. T., Wedge, D. C., Van Loo, P., PCAWG Evolution and Heterogeneity Working Group, and PCAWG Network (2017). The evolutionary history of 2,658 cancers. bioRxiv, page 161562.

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## Consortium authorships

Abbosh, C.\*, Birkbak, N. J.\*, Wilson, G. A.\*, Jamal-Hanjani, M.\*, Constantin, T.\*, Salari, R.\*, Quesne, J. L., Moore, D. A., Veeriah, S., Rosenthal, R., Maroti, T., Kirkizlar, E., Watkins, T. B. K., McGranahan, N., Ward, S., Martinson, L., Riley, J., Fraioli, F., Bakir, M. A., Gronroos, E., Zambrana, F., Endozo, R., Bi, W. L., Fennessy, F. M., Sporer, N., Johnson, D., Laycock, J., Sha, S., Czyzewska-Khan, J., Rowan, A., Chambers, T., Matthews, N., Turajlic, S., Hiley, C., Lee, S. M., Forster, M. D., Ahmad, T., Falzon, M., Borg, E., Lawrence, D., Hayward, M., Kolvekar, S., Panagiotopoulos, N., Janes, S. M., Thakrar, R., Ahmed, A., Blackhall, F., Summers, Y., Hafez, D., Naik, A., Ganguly, A., Kareht, S., Shah, R., Joseph, L., Quinn, A. M., Crosbie, P., Naidu, B., Middleton, G., Langman, G., Trotter, S., Nicolson, M., Remmen, H., Kerr, K., Chetty, M., Gomersall, L., Fennell, D. A., Nakas, A., Rathinam,

S., Anand, G., Khan, S., Russell, P., Ezhil, V., Ismail, B., Irvin-sellers, M., Prakash, V., Lester, J. F., Kornaszewska, M., Attanoos, R., Adams, H., Davies, H., Oukrif, D., Akarca, A. U., Hartley, J. A., Lowe, H. L., Lock, S., Iles, N., Bell, H., Ngai, Y., Elgar, G., Szallasi, Z., Schwarz, R. F., Herrero, J., Stewart, A., Quezada, S. A., Van Loo, P., Dive, C., Lin, C. J., Rabinowitz, M., Aerts, H. J., Hackshaw, A., Shaw, J. A., Zimmermann, B. G., the TRACERx Consortium, the PEACE Consortium, and Swanton, C. (2017). Phylogenetic ctDNA analysis depicts early stage lung cancer evolution. *Nature*.

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