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Connectivity of diagnostic technologies: improving surveillance and accelerating TB elimination

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<http://www.ingentaconnect.com/content/iuatld/ijtld/2016/00000020/00000008/art00004>

Authors

Emmanuel ANDRE

- Pôle de Microbiologie, Institut de Recherche Expérimentale et Clinique, Université catholique de Louvain, Belgium
- Service de Microbiologie, Département de Biologie Clinique, Cliniques Universitaires Saint-Luc, Belgium
- ESGMYC Study Group for Mycobacterial Infections, European Society for Clinical Microbiology and Infectious Diseases

Chris ISAACS

- Foundation for Innovative New Diagnostics, Switzerland

Dissou AFFOLABI

- Faculty of Health Sciences, Abomey-Calavi University, Benin
- National Tuberculosis Programme, Benin

Riccardo ALAGNA

- TB Supranational Reference Laboratory, IRCCS San Raffaele Scientific Institute, Italy

Dirk BROCKMANN

- Institute for Theoretical Biology, Department of Biology, Humboldt University of Berlin, Germany
- Epidemiological Modelling of Infectious diseases, Robert Koch-Institute, Germany

Bouke C de JONG

- 28 • Unit of Mycobacteriology, Department of Biomedical Sciences, Institute of Tropical Medicine,
29 Belgium
- 30 Emmanuelle CAMBAU
- 31 • Université Paris Diderot, INSERM UMR 1137 IAME, France
- 32 • APHP, Hôpital Lariboisière, Bactériologie, France
- 33 • ESGMYC Study Group for Mycobacterial Infections, European Society for Clinical Microbiology and
34 Infectious Diseases
- 35 Gavin CHURCHYARD
- 36 • Aurum Institute, South-Africa
- 37 Ted COHEN
- 38 • Yale School of Public Health, United States of America
- 39 Michel DELMEE
- 40 • Pôle de Microbiologie, Institut de Recherche Expérimentale et Clinique, Université catholique de
41 Louvain, Belgium
- 42 • Service de Microbiologie, Département de Biologie Clinique, Cliniques Universitaires Saint-Luc,
43 Belgium
- 44 Jean-Charles DELVENNE
- 45 • Institute of Information and Communication Technologies, Electronics and Applied Mathematics,
46 Université catholique de Louvain, Belgium
- 47 • Centre for Operations Research and Econometrics, Université catholique de Louvain, Belgium
- 48 Maha FARHAT
- 49 • Massachusetts General Hospital, United States of America
- 50 Ali HABIB
- 51 • Interactive Health Solutions, Pakistan
- 52 Petter HOLME
- 53 • Sungkyunkwan University, South Korea
- 54 Salmaan KESHAVJEE
- 55 • Harvard Medical School Center for Global Health Delivery–Dubai, United Arab Emirates

- 56 Aamir KHAN
- 57 • Interactive Research and Development, Pakistan
- 58 Piedra LIGHTFOOT
- 59 • Foundation for Innovative New Diagnostics, Switzerland
- 60 David MOORE
- 61 • TB Centre, London School of Hygiene and Tropical Medicine, United Kingdom
- 62 Yamir MORENO
- 63 • Institute for Biocomputation and Physics of Complex Systems (BIFI), University of Zaragoza, Spain
- 64 • Department of Theoretical Physics, Faculty of Sciences, University of Zaragoza, Spain
- 65 Yamuna MUNDADE
- 66 • UNITAID, Switzerland
- 67 Madhukar PAI
- 68 • McGill International TB Centre & McGill Global Health Programs, McGill University, Canada
- 69 Sanjay PATEL
- 70 • University Hospital Southampton NHS Foundation Trust, United Kingdom
- 71 Alaine Umubyeyi NYARUHIRIRA
- 72 • Management Sciences for Health, South Africa
- 73 Luis E C ROCHA
- 74 • Karolinska Institutet, Sweden
- 75 • Université de Namur, Belgium
- 76 Jeff TAKLE
- 77 • Global Connectivity LLC, United States of America
- 78 Arnaud TREBUCQ
- 79 • International Union against Tuberculosis and Lung Disease, France
- 80 Jacob CRESWELL
- 81 • Stop TB Partnership, Switzerland
- 82 Catharina BOEHME
- 83 • Foundation for Innovative New Diagnostics, Switzerland

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87 **ABSTRACT IN ENGLISH**

88 In regard to tuberculosis (TB) and other major global epidemics, the use of new diagnostic tests is
89 increasing dramatically, including in resource-limited countries. Although there has never been as much
90 digital information generated, this data source has not been exploited to its full potential. In this opinion
91 paper, we discuss lessons learned from the global scale-up of these laboratory devices and the pathway
92 to tapping the potential of laboratory-generated information in the field of TB by using connectivity.
93 Responding to the demand for connectivity, innovative third-party players proposed solutions that have
94 been widely adopted by field users of the Xpert MTB/RIF assay. The experience associated with the
95 utilization of these systems, which facilitate the monitoring of wide laboratory networks, stressed the
96 need for a more global and comprehensive approach to diagnostic connectivity. In addition to
97 facilitating the reporting of test results, the mobility of digital information allows the sharing of
98 information generated in programme settings. These data, when they become easily accessible, can be
99 used to improve patient care, disease surveillance and drug discovery. Therefore they should be
100 considered as a public health good. We list several examples of concrete initiatives that should allow
101 data sources to be combined to improve the understanding of the epidemic, support the operational
102 response, and finally accelerate TB elimination. With the many opportunities that the pooling of data
103 associated with the TB epidemic can provide, pooling of this information at an international level has
104 become an absolute priority.

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109 **Résumé en Français**

110 Dans le domaine de la tuberculose (TB) et d'autres épidémies majeures au niveau international,
111 l'utilisation de nouvelles technologies pour le diagnostic s'est largement répandue, y compris dans les
112 pays à faible ressources. Cependant, malgré la grande quantité de données générées par ces nouveaux
113 outils, cette source d'information reste aujourd'hui largement inexploitée. Dans cet article d'opinion,
114 nous discutons les leçons tirées de l'utilisation de ces nouveaux outils diagnostics et certaines pistes
115 pour mieux mettre à profit, grâce à la connectivité, les informations générées par les laboratoires TB. En
116 réponse à l'absence de solutions permettant cette connectivité, des solutions innovantes ont été
117 proposées par des acteurs tiers et ont été largement adoptées par les utilisateurs du test Xpert
118 MTB/RIF. L'utilisation croissante de ces solutions qui permettent la surveillance de larges réseaux de
119 laboratoires a porté l'attention sur la nécessité de proposer une approche plus globale et intégrée par
120 rapport à la connectivité des laboratoires diagnostiques. Ces solutions facilitent la transmission des
121 résultats, mais permettent également le partage d'informations générées en situation réelle. Ces
122 données, lorsqu'elles deviennent aisément accessibles, peuvent être utilisées pour améliorer la qualité
123 des soins prodigués aux malades, la surveillance des maladies et la découverte de médicaments. Pour
124 ces raisons, elles doivent être considérées comme un bien de santé publique. Nous dressons une liste
125 d'exemples d'initiatives concrètes qui devraient permettre de faciliter le partage de données de
126 laboratoire dans le but de renforcer notre compréhension de l'épidémie, soutenir les réponses
127 opérationnelles, et accélérer l'élimination de la TB. En raison des nombreuses opportunités associées au
128 partage d'information liées à l'épidémie de TB, la centralisation des données au niveau international est
129 devenue une priorité absolue.

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132 **Resumen en español**

133 En el contexto de la tuberculosis (TB), la utilización de nuevas pruebas diagnósticas está aumentando de
134 manera espectacular, especialmente en los países en desarrollo. Pese a que nunca se ha generado tanta
135 cantidad de datos, aún no se aprovechan todas las posibilidades que ofrece esta nueva fuente de
136 información. En el presente artículo de opinión, se examinan las enseñanzas extraídas del uso en todo el
137 mundo de estos nuevos instrumentos diagnósticos y se analiza la hoja de ruta hacia la explotación de las
138 ventajas y el potencial de la conectividad para el diagnóstico de la TB. Respondiendo a la falta de
139 conectividad incorporada a las herramientas de diagnóstico, se han creado soluciones de conectividad,
140 que a su vez han sido adoptadas por usuarios en el terreno con el fin de monitorizar la utilización del
141 test Xpert MTB/RIF. El uso creciente de estas soluciones ha centrado la atención sobre la necesidad de
142 explorar de manera más general y exhaustiva la conectividad destinada al diagnóstico. Además de
143 facilitar a los laboratorios la tarea de comunicar los resultados, la información digital debería favorecer
144 el intercambio y el acopio de la información recogida en el marco programático. Dado que estos datos
145 pueden mejorar la atención al paciente, la vigilancia de enfermedades y el descubrimiento de nuevos
146 medicamentos, es preciso considerarlos como un bien de salud pública. Aquí, enumeramos varios
147 ejemplos de iniciativas concretas que deberían facilitar la combinación de diferentes fuentes de datos
148 para mejorar la vigilancia de la TB y acelerar su eliminación. Habida cuenta de las múltiples soluciones
149 que ofrece, la combinación de datos a escala internacional constituye una prioridad absoluta, pues
150 agilizará el progreso en sectores primordiales como la atención al paciente, la vigilancia epidemiológica
151 y la respuesta operativa.

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153 **Background**

154 In the past decade, the use of new diagnostic tests has increased dramatically in developing
155 countries' laboratories and more recently in decentralised point-of-care facilities. Self-contained
156 molecular diagnostic devices have been successfully deployed to detect tuberculosis (TB) (e.g.
157 GeneXpert¹) or monitor treatment for HIV (e.g. PIMA²) in very basic clinical facilities.

158 Despite the accumulating evidence that these tools can be successfully used in the most challenging
159 environments^{3,4} and the establishment of distribution and funding channels that should theoretically
160 allow any country to access and scale-up these new technologies, the majority of patients that could
161 benefit from these technical evolutions still do not have access to them. It is clear that the
162 introduction of an improved TB diagnostic is not sufficient for assuring improved outcomes for
163 patients as the details of implementation within existing health-delivery systems have critical
164 influence on impact⁵. We suggest that the introduction of new tools such as GeneXpert offers an
165 important opportunity to better understand, monitor and improve such delivery systems to assure
166 greatest impact. If scale-up of novel diagnostic devices can be accompanied by the simultaneous
167 introduction of up-to-date quality indicators and technical connectivity solutions, the vast amount of
168 data generated by these new generation of automates could actually both simplify and potentiate
169 the global response to the TB epidemic.

170 On a national and global level, the quantity of information produced following the introduction of
171 new-generation laboratory instruments was not anticipated, thus there were no plans in place for
172 how to manage the information flow or orient it in such a way that it could generate an evolution in
173 the organisation of the epidemic response. In the absence of adequate laboratory information
174 technology infrastructure, complemented with standardised reporting solutions for screening
175 activities and treatment follow-up, many low-resource countries have continued to use slow and
176 error-prone paper-based recording systems. In such systems, editing and transmission of paper
177 reports cause inherent delays and contribute to the cost, complexity and relative inaccuracy of data

178 interpretation.

179 Diagnostic ehealth solutions have the potential to help overcome some of these problems and
180 maximize patient and public health impact following the introduction of a particular technology. The
181 combination of this unprecedented evolution of the laboratory landscape and the potential of
182 eHealth could be leveraged to generate evolution in national and global health-delivery systems that
183 is needed to achieve TB elimination. Pragmatically, this requires device connectivity, wherein testing
184 data and results are automatically and securely sent to repositories, translated into useful
185 information and channeled to appropriate parties. Although device connectivity within other
186 industries has been commonplace for some time, within the healthcare community it is still
187 considered to be in its infancy ⁶.

188 In this paper, we discuss lessons learned from the global scale-up of the first generation of easy-to-
189 connect diagnostic tools ⁷ and the pathway to tapping the potential of connectivity in the field of TB
190 diagnostics ⁸.

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192 **Experience from first-generation connected diagnostics: the example of Xpert MTB/RIF**

193 During the last decade, several diagnostic companies, such as Cepheid Inc. (Sunnyvale, USA) and Alere
194 Inc. (Waltham, USA), began developing a new generation of tests—essential to fight diseases of poverty
195 such as TB and HIV—with significant support from public and philanthropic funders including NIH and
196 BMGF.

197 The Xpert MTB/RIF, run on the GeneXpert platform, was the first truly game-changing test to come out
198 of this work and has since been widely distributed in health facilities with limited human and
199 infrastructure resources. The coverage of GeneXpert varies importantly between countries, with some
200 countries still having only a couple of machines based in reference laboratories, and other countries
201 such as South-Africa which rapidly realized the advantages of implementing this novel platform as a

202 first-line test⁹. In the last five years, more than 13 million Xpert MTB/RIF tests have been procured
203 worldwide. When GeneXpert was rolled out in 2010, the instrument had no built-in connectivity outside
204 basic standards and the TB community did not have the software tools to connect to GeneXpert
205 machines and optimally use the data being generated. As a consequence, valuable information was
206 housed in the hard drives of local computers, was never used to inform surveillance efforts or health
207 care providers, and has largely been lost.

208 In light of this, national TB programmes called for tools to reduce loss to follow-up and improve device
209 and laboratory management—including a better ability to maintain cartridge supply and local
210 redistribution and to evaluate and fulfill the training needs of device operators and lab technicians.
211 Likewise, TB programmes voiced a need for connectivity systems that could relieve the high overhead
212 costs of data aggregation and analysis that hamstrings the process of collecting raw data and turning it
213 into useful information.

214 In 2012, responding to this critical gap in the implementation landscape, innovative third-party players
215 developed connectivity solutions. For example, GxAlert (ABT and SystemOne), XpertSMS (Interactive
216 Research and Development and TB REACH) and GenXchange (Université catholique de Louvain and the
217 National TB Program of DRC) were devised to respond to the needs of low-resource countries where
218 often internet is unavailable or unreliable and laboratory information systems or electronic medical
219 records are not widely used. These tools offered immediate solutions and, based on national requests,
220 hundreds of local laboratories have since been interconnected by implementing these systems. The
221 scaling of these connectivity solutions has been taken back by dedicated companies^{10,11}.

222 Cepheid, the manufacturer of GeneXpert, also worked to enable remote monitoring of their devices in
223 response to expressed national needs and requests from the TB community. Like many developers,
224 Cepheid lacked comprehensive information about what use-cases needed to be supported, and for
225 ethical and regulatory reasons prioritized data security and confidentiality. As a result, the company
226 launched an initial software tool that was a step forward but unable to fulfill all programme needs.

227 In response, a WHO-led alliance of key implementation partners (e.g. USAID, MSF, CHAI and FIND) and
228 donors (e.g. UNITAID and GFATM) was formed to work with Cepheid to ensure secure, open access to
229 critical data and to find a broader, holistic approach to connectivity and data management. An
230 immediate solution was found and both Cepheid and the alliance remain interested in the creation of a
231 non-proprietary, long-term connectivity platform (or a series of integrated and inter-operational
232 platforms). This highlights how the global TB community can collectively define priority needs and work
233 with manufacturers to negotiate and realize solutions for accessing and utilising key data.

234 Another important lesson from the implementation of first-generation connected diagnostics is the
235 importance of a well-tailored delivery pathway for connectivity software that supports sustainable up-
236 take in country. For instance, Alere, the manufacturer of PIMA, devised a country-based public-private
237 partnership model to ensure appropriate training and support for their connectivity software. Without
238 this support and engagement of key stakeholders, many countries would have struggled to make use of
239 the influx of data. While the tool itself has limited wider applicability because of the proprietary nature
240 of the software, the partnership model offers a valuable example of how non-proprietary, interoperable
241 systems could be disseminated and nurtured in the future.

242 **Connectivity of diagnostics: a shared responsibility and public health necessity**

243 WHO and research funding agencies have been advocating for, and implementing, data-sharing policies
244 for some time. While these efforts have increased access to synthesized research data, efforts to make
245 national programme data available are in their infancy. The use of new generation diagnostic platforms
246 has triggered thinking about the potential utility of real-time analysis of national data and how
247 diagnostic connectivity could further improve epidemiological surveillance and guide targeted public
248 health responses. Accelerated TB elimination, for example, as called for in the WHO End TB strategy ¹²,
249 can only be realized if case detection, individual patient management and epidemiological surveillance
250 are intensified simultaneously, and if these efforts are closely monitored and validated. Data generated
251 by Xpert MTB/RIF testing can be used both to improve patient management and treatment efforts, and

252 to provide important population-level information on average infectiousness as a predictor for TB
253 burden¹³ and spread of new mutations. This requires optimized programmatic data management,
254 pooling, sharing, analysis and use. Realizing improvements in surveillance and public health demands
255 that information generated by diagnostic technologies in programmatic conditions be easily accessible
256 and usable for national programmes. Ultimately, data access, enabled by diagnostic connectivity, should
257 thus be seen as a public health good. Countries, international organizations, test developers and civil
258 society organizations have a collective responsibility to work together to ensure sustainable use of
259 information and communications technology to improve healthcare. In doing so, important questions
260 regarding ethical obligations, data ownership and stakeholder interests, e.g. market competitiveness,
261 need to be acknowledged and addressed. International collaborative efforts must furthermore address
262 the issue of personal unique identifiers in a context of continuous human migrations and data mobility.

263 **The way forward: realizing the potential of connected diagnostics**

264 Built-in connectivity has become an evident prerequisite for upcoming diagnostic platforms¹⁴. Tests that
265 until recently were un-connectable, such as rapid diagnostic tests (e.g. HIV, malaria), can now be
266 connected to digital readers with results collected, stored and transferred (e.g. Fio Corp, Canada).

267 In the field of TB diagnostics, a wide range of laboratory tests are used in complementarity. This includes
268 rapid diagnostic tests and more conventional approaches such as microscopy, culture, drug
269 susceptibility testing and sequencing¹⁵. Inter-connecting these diagnostic devices and further
270 integrating this information with clinical indicators is the upcoming challenge for the TB community.

271 The Connected Diagnostics Initiative (CDx), coordinated by FIND (Geneva, Switzerland), is an example of
272 a potential solution for accelerating connectivity and interoperability of diagnostic devices. CDx is
273 providing an open-source software platform allowing for centralised aggregation of data from
274 diagnostics, regardless of manufacturer. For this new effort to succeed, wide buy-in from implementers,
275 policymakers and developers will be essential. In parallel, FIND is working with WHO towards guidelines
276 for standardised results reporting for diagnostic devices, and assisting developers to be in compliance

277 with these standards. These efforts go hand in hand with further deployment of local laboratory
278 information systems and electronic medical records ¹⁶.

279 Alongside this initiative, various groups are creating global databases with the intention of enhancing
280 research and development applications of data. For instance, genTB (Harvard University) is an open-
281 source platform that allows for the pooling, analysis and visualization of genetic, epidemiological and
282 clinical data. A global partnership, including WHO, CDC, CPATH, Stop TB, NIAID and FIND, has been
283 established to develop a data platform (ReSeqTB) to store, curate and provide access to globally
284 representative TB data that can inform the development of new diagnostics, facilitate clinical decisions
285 and improve surveillance of drug resistance. While the opportunities for sharing information at an
286 international scale must be promoted, countries must be provided with technical solutions that can
287 support them in efficiently managing with whom, and for what purposes, national data are shared, and
288 to ensure that these database efforts ultimately benefit patients.

289 Consensus is forming around the central role that connected diagnostics and digitization can play in
290 tackling health systems weaknesses and diseases of poverty. However, the global health community
291 must also address complex question of how new tools and practices can be effectively implemented in
292 health systems. Substantial programmatic changes will be required in countries to absorb the innovation
293 of connectivity and capture its benefits. This demands a holistic approach to cultivating effective
294 development and adoption of new diagnostic tools. In this context, laboratory connectivity may serve
295 the need for more efficient post-marketing surveillance of newly rolled-out diagnostics both for national
296 stakeholders and their global partners. As the amount of information collected will rapidly increase
297 beyond our conventional capacities of analysis, the global health community will also need to initiate
298 and intensify innovative collaborations to exploit the data collected, using big data analysis and self-
299 learning algorithms. Managing, visualizing and analysing such big data creates challenges beyond the
300 capacities of standard statistical methods, and thus generates an increasing demand for data science
301 and multidisciplinary efforts.

302 **Conclusion**

303 Our common goal of TB elimination is not a dream anymore: it is an achievable objective with clear
304 milestones ¹⁷. The elimination effort will require a strengthened collaboration between information
305 technology and big data specialists, social medicine and private companies ⁶.

306 In the future, all diagnostic technologies should be inter-connected, allowing data generated by
307 laboratories to merge in a common repository while safeguarding patient confidentiality. The TB
308 community could use such a repository to monitor progress and identify problems and potential
309 solutions, at both patient and global levels. Data pooling will open up opportunities to comprehend the
310 rapid evolution of drug-resistant mutations, which will aid in selecting cost-efficient treatment schemes
311 and improving patient management. With the many solutions it can provide, data pooling at an
312 international level is an absolute priority, as it will accelerate progress in critical sectors including patient
313 care, epidemiological surveillance and operational response. Being an international health emergency,
314 the TB epidemic requires optimal international collaboration and unambiguous political commitment for
315 intensifying data sharing efforts.

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