

Data collection for machine design of novel antibiotics against ESKAPE bacteria

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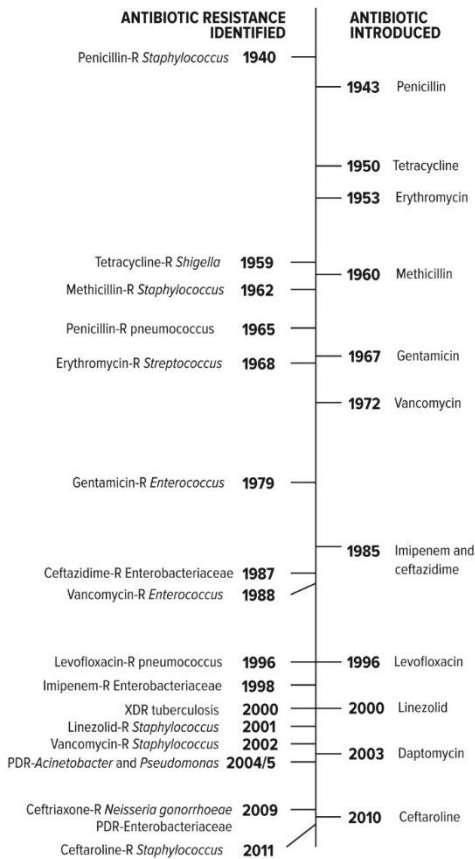
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Antibiotic Resistance Crisis

In 2019 more than 2.8 million antibiotic-resistant infections occurred in the U.S., and more than 35,000 people died

**Figure 1 Developing Antibiotic Resistance:
A Timeline of Key Events⁵**



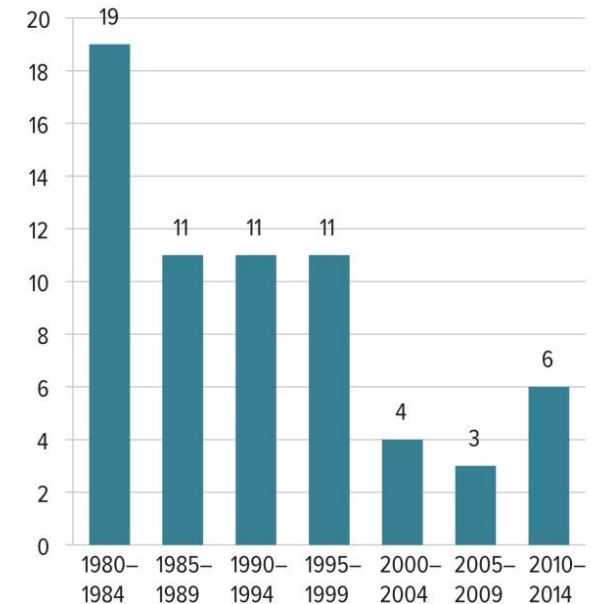
PDR = pan-drug-resistant; R = resistant; XDR = extensively drug-resistant

Dates are based upon early reports of resistance in the literature. In the case of pan-drug-resistant *Acinetobacter* and *Pseudomonas*, the date is based upon reports of health care transmission or outbreaks. Note: penicillin was in limited use prior to widespread population usage in 1943.

1. Old antibiotics are no longer working.
2. Antibiotics are no longer favorable investments.

- a. Expensive to research.
- b. Doctors are less likely to prescribe new antibiotics .
- c. New antibiotics are dedicated to only treating severe infections or resistant infections.

Figure 3 Number of Antibacterial New Drug Application Approvals Versus Year Intervals



The number of new antibiotics developed and approved has decreased steadily over the past three decades (although four new drugs were approved in 2014), leaving fewer options to treat resistant bacteria.

* Drugs are limited to systemic agents. Data courtesy of the CDC⁵ and the FDA Center for Drug Evaluation and Research.

from C. Lee Ventola *PT*, 2015, 40, 277



ESKAPE Bacteria -Six Highly Antibiotic Resistance Bacterial Pathogens

Enterococcus faecium - gram-positive bacteria

Staphylococcus aureus - gram-positive bacteria

Klebsilla pneumoniae - gram-negative bacteria

Acinetobacter baumannii - gram-negative bacteria

Pseudomonas aeruginosa - gram-negative bacteria

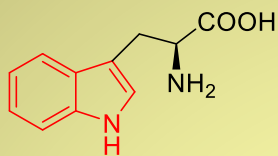
Enterobacter - gram-negative bacteria

- **700,000 deaths per year are estimated to be caused by gram- negative bacteria world wide.**
- **Most hospital acquired infections are due to ESKAPE bacteria.**

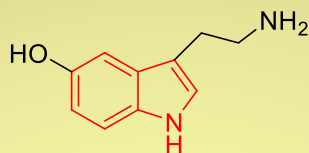
Koulenti, D., et al. (2018). "Infections by multidrug-resistant Gram-negative Bacteria: what's new in our arsenal and what's in the pipeline?"; Peleg, A. Y. , D. C. Hooper, *N. Engl. J. Med*, **2010**, 362, 1804.



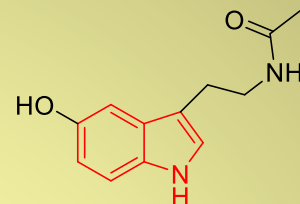
Examples of Natural Compounds and Drugs Containing Indole Core



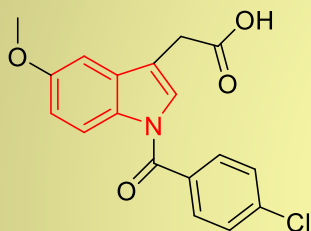
Tryptophan



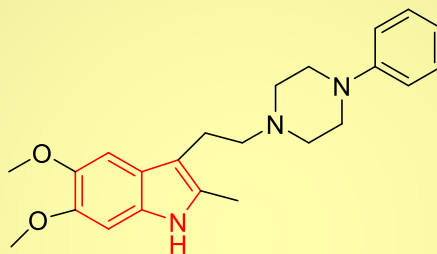
Serotonin



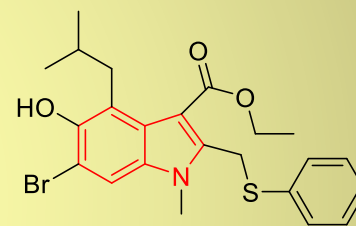
Melatonin



Indometacin



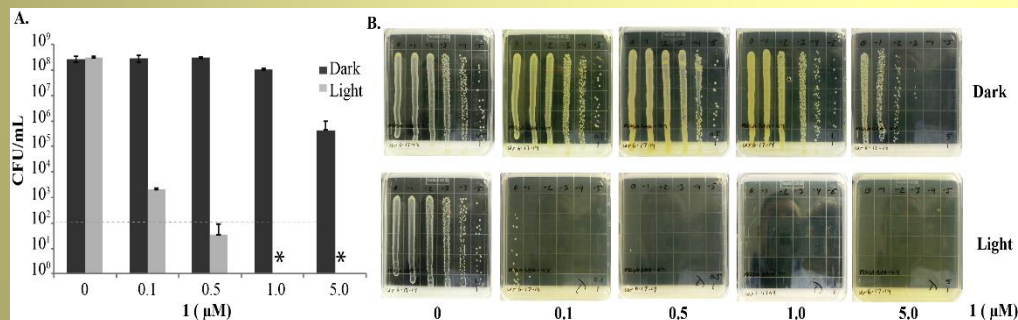
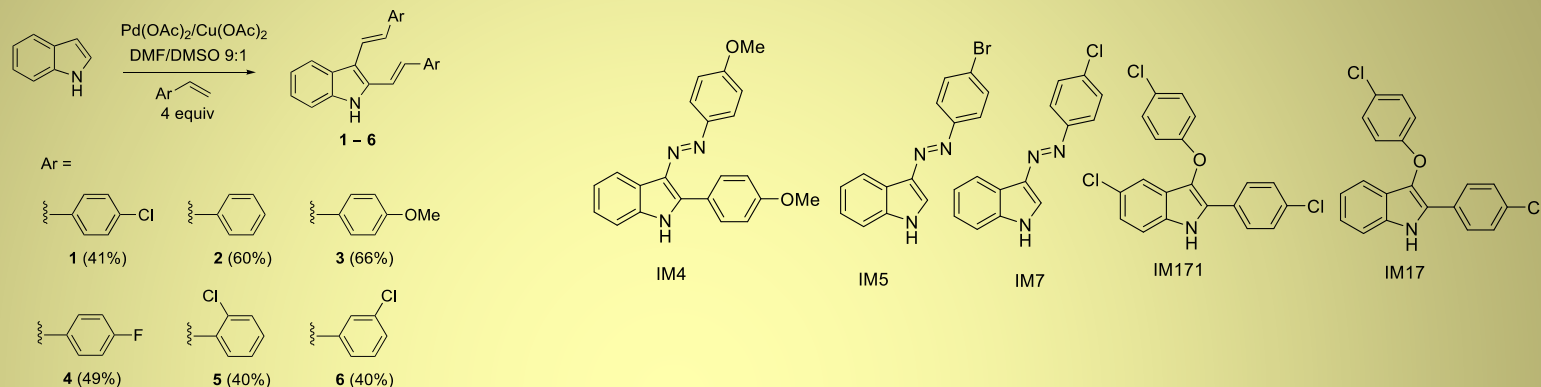
Oxypertine



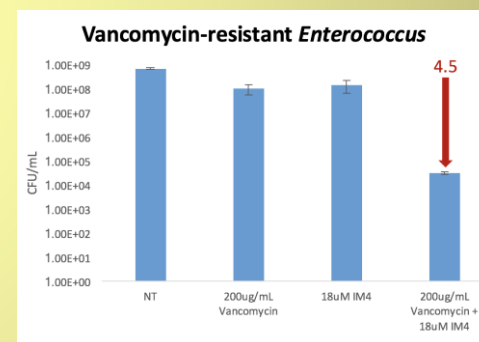
Arbidol

Drug	Application	Drug	Application
Vincristine	Anticancer	Vincamine	Vasodilator
Vinblastine	Anticancer	Roxindole	Schizophrenia
Reserpine	Antihypertensive	Delavirdine	Anti-HIV
Vinorelbine	Anticancer	Peridopril	Antihypertensive
Vindesine	Anticancer	Binedaline	Antidepressant
Atevirdine	Anti-HIV	Zafirlukast	Anti-Asthmatic
Mitraphylline	Anticancer	Amedalin	Antidepressant
Cediranib	Anticancer	Oxypertine	Antipsychotic

New Antibacterial Compounds Against Resistant Bacteria



Compound 1 eradicates MRSA (ATCC BAA 44) in the presence of light. A: Graph, showing a dose-dependent effect of 1 with and without a 2-minute irradiation with white light. B: Drip-streak plates utilized in the experiment illustrating the CFU counts. The data are derived from a single experiment performed in 3 replicates and serve as an example of 3 independent experiments producing similar results. The stars represent no observable colonies.



The antibacterial activities were determined using an optimized time kill method. The error bars represent technical triplicates in a representative biological replicate carried out at least three times



Design and Development of Next Generation Antibiotics Using Machine Learning Approach

Target identification and validation	Compound screening and lead discovery	Preclinical development
<ul style="list-style-type: none"> • Target identification and prioritization based on gene–disease associations • Target druggability predictions • Identification of alternative targets (splice variants) 	<ul style="list-style-type: none"> • Compound design with desirable properties • Compound synthesis reaction plans • Ligand-based compound screening 	<ul style="list-style-type: none"> • Tissue-specific biomarker identification • Classification of cancer drug–response signatures • Prediction of biomarkers of clinical end points
<ul style="list-style-type: none"> • Current data are highly heterogeneous: need standardized high-dimensional target–disease–drug association data sets • Comprehensive omics data from disease and normal states • High-confidence associations from the literature • Metadata from successful and failed clinical trials 	<ul style="list-style-type: none"> • Large amounts of training data needed • Models for compound reaction space and rules • Gold standard ADME data • Numerous protein structures 	<ul style="list-style-type: none"> • Biomarkers: reproducibility of models based on gene expression data • Dimension reduction of single-cell data for cell type and biomarker identification • Proteomic and transcriptomic data of high quality and quantity

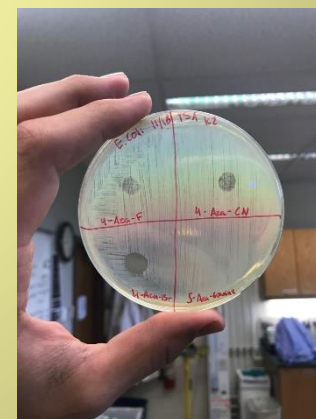
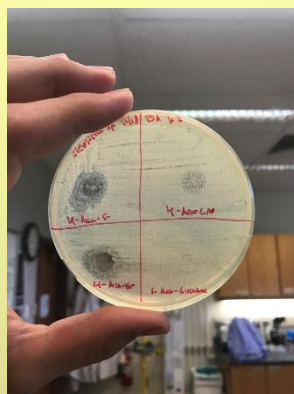
from J. Vamathevan *Nat. Rev. Drug. Discov.*, **2019**, 18, 463



First Results

Drug-like parameters and biological data were collected for about 40 indole derivatives.
A first library of novel potential antibiotics was synthesized.

Strain of bacteria	Name of new compound					
	4-Aza-H	5-Aza-CN	5-Aza-Br	7-Aza-Gu	7-Aza-F	7-Aza-Br
<i>E. coli</i>	+	-	+	-	-	-
<i>E. carotovora</i>	+	+	+	+	+	-
<i>Klebsiella</i> sp.	+	-	+	-	-	-
<i>S. epidermidis</i>	-	-	+	-	-	-
<i>P. aeruginosa</i>	+	-	+	+	+	+
<i>B. subtilis</i>	+	-	+	-	-	-



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