



# Implications of access and benefit-sharing (ABS) frameworks for collection and utilisation of marine biological samples

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# Why Use Marine Bioresources?

Offers advantage over comparable terrestrial resource:

- Superior performance

- Better economics

Unprecedented activity in particular application:

- Enzymes: new reactivity/new biotransformation

- Small molecules: novel chemical structures & new mechanism of action

- Materials: new properties

# Marine Biotechnology Products on the Market



Vent Polymerase

Origin: Vent bacterium

Production: Recombinant



Priali for pain

Origin: Phillippino cone snail

Production: Recombinant



$\omega$ -3 polyunsaturated fatty acids

for heart disease Source: Fish

Production: Fish



Halaven for cancer

Origin: Japanese deep water sponge

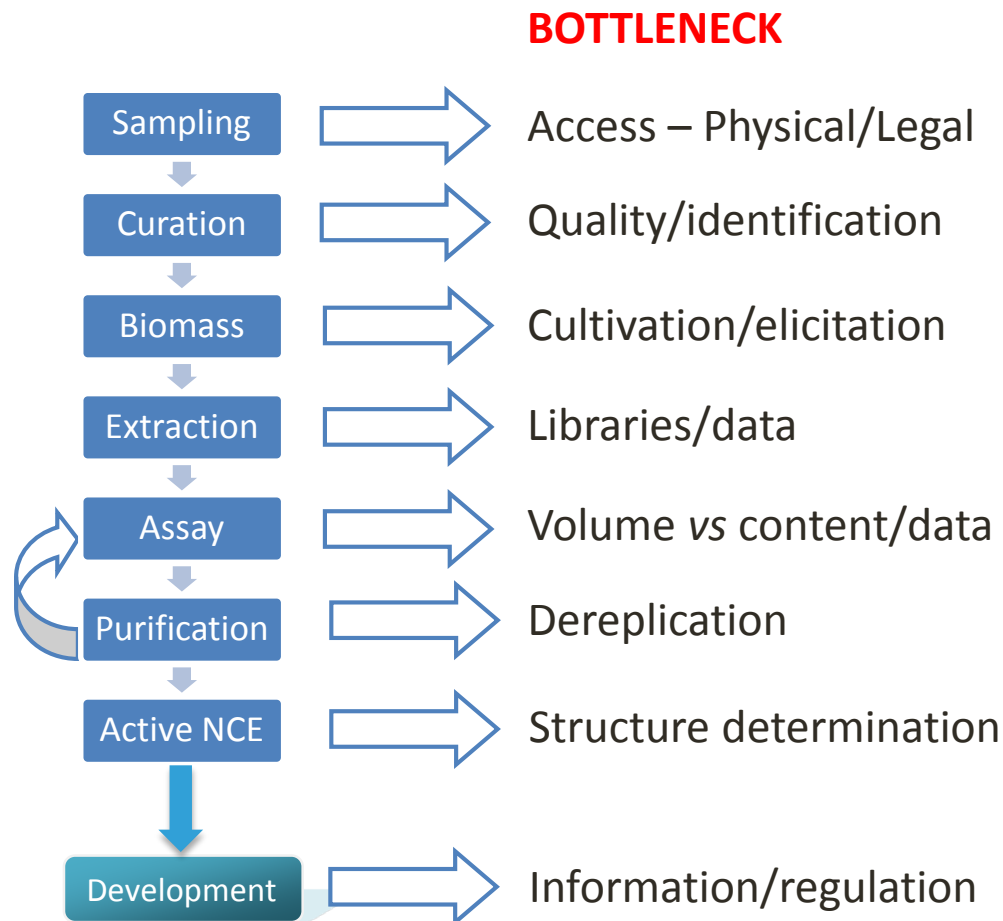
Production: Chemical synthesis

# The Marine Biodiscovery Process

**Biodiscovery** is the discovery of compounds and associated ideas from natural sources to develop novel biomedicines.

**Biodiscovery** generates chemical diversity that is used to find initial biological activity in disease focused screens

**Biodiscovery** also includes the development of biomedical research tools, antifoulants, catalysts, nutraceuticals and cosmeceuticals.



# Applying ABS Nagoya-Style to BBNJ

- ABNJ does not refer to any ecological habitat or environment
- MGR has no real meaning to scientists
- May inhibit research due to difficulties in gaining access
  - Where stringent agreements exist few commercial deals have been done (Australia)
- Will it require an equivalent to access with prior informed consent?
- How will it be monitored/policed?
- Who will collect monetary benefits (ISA?) and who will distribute funds and how? How will terms be agreed (no MAT)?
- Becomes multilateral relationship not bilateral as in NP.
- Traceability becomes an issue as benefits may take a long time to be realised (& who will do this?).
- Expectations of financial returns may be very unrealistic.

# Real Benefit Scenario

- Cost in 2014 to bring drug to market US\$2,558 M\* - >70% Clinical trials
- Typical industry royalties on natural products developed into drugs is 1-3%
- Halaven (Eisai), derived from a Japanese sponge makes US\$200 M per year – in principle yielding US\$ 2-6 M pa.
- Currently 7 approved marine drugs – total royalties would be US\$ 10-50 M.
- Blockbuster drug (> US\$ 1 Bn pa income) would yield US\$10-30 M pa
- Currently 7 approved marine drugs come from ~28,000 discovered marine compounds (1 in 4000 chance) – none are ‘blockbusters’
- All examples were discovered pre-CBD – not clear if actual royalties are being paid

\*Tufts Study [http://csdd.tufts.edu/news/complete\\_story/cost\\_study\\_press\\_event\\_webcast](http://csdd.tufts.edu/news/complete_story/cost_study_press_event_webcast)

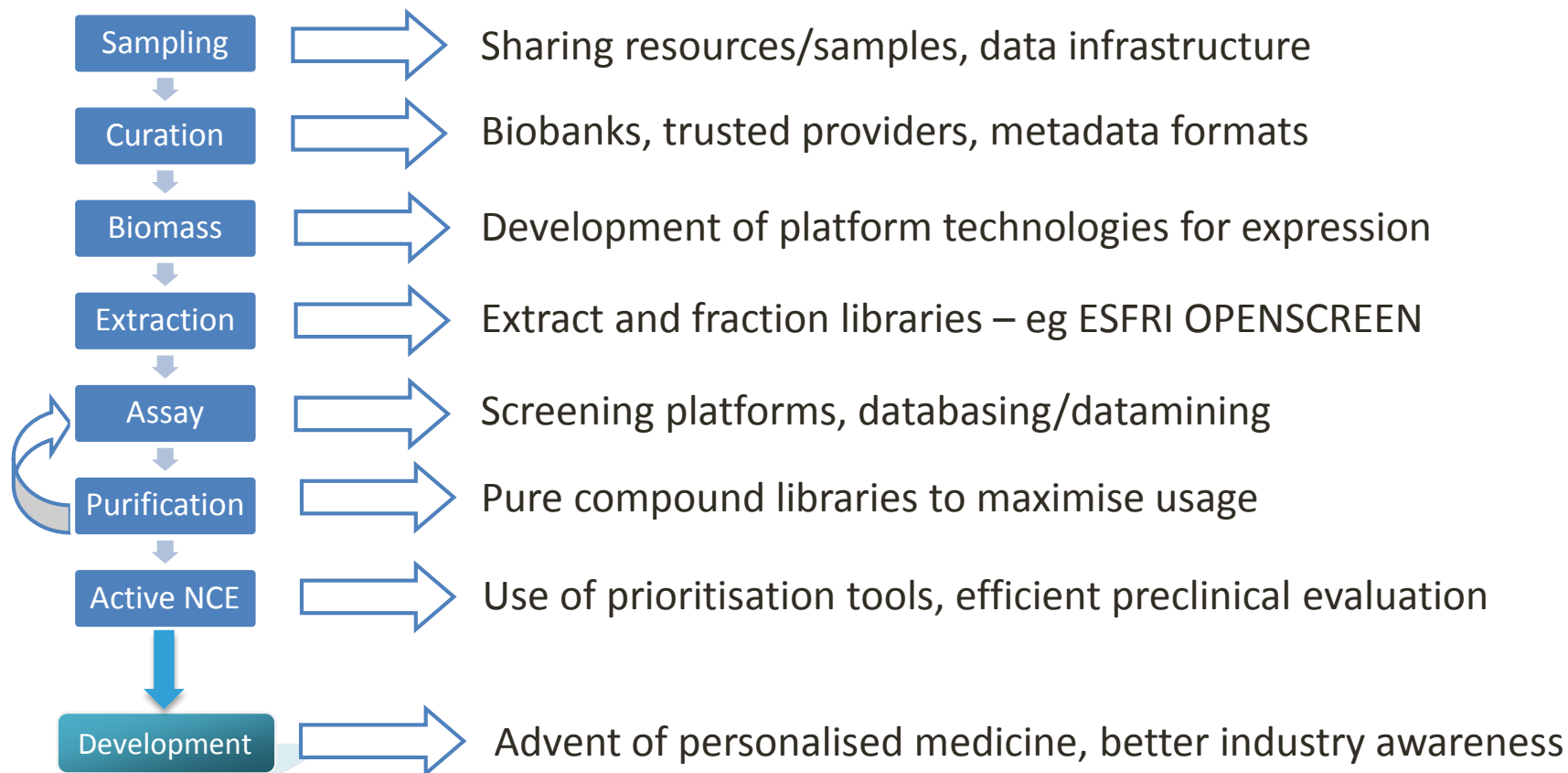
# The Major Benefits are Non-Monetary

- In many cases most important benefits from use of bioresources
- In many cases these are the only benefits
- Non-monetary benefits may include:
  - Scientific exchanges/training
  - Technology transfer
  - Capacity building (infrastructure)
  - Enhanced reputation
  - Increased number/quality of scientific publications
  - Biodiversity conservation
  - Valuable regional resources developed (knowledge, samples, data)
- Non-monetary benefits still cost money – however they are upfront compared to royalties
- Is a Nagoya-like regime necessary or desirable? Will good practice suffice?

# The Marine Biodiscovery Process

## Elements of Good Practice

### Current Good Practice





# Current Good Practice in Cruise Planning

## Application

- Cruise path/stations/equipment.

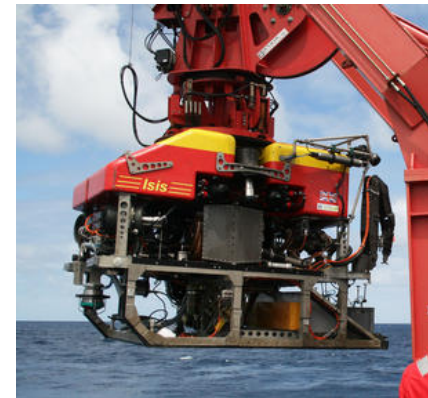


## Award

- Clarification for feasibility/equipment availability.
- Check MPAs not entered.

## After Cruise

- Data is logged with central agency – cruise report
- Sample list/locations collected/location stored
- Environmental data/images and video



**Currently no requirement for post cruise data (eg genetic data) to be deposited. Species may not be identified until later, if at all**  
**Responsible sampling protocols developed (eg Interridge)**

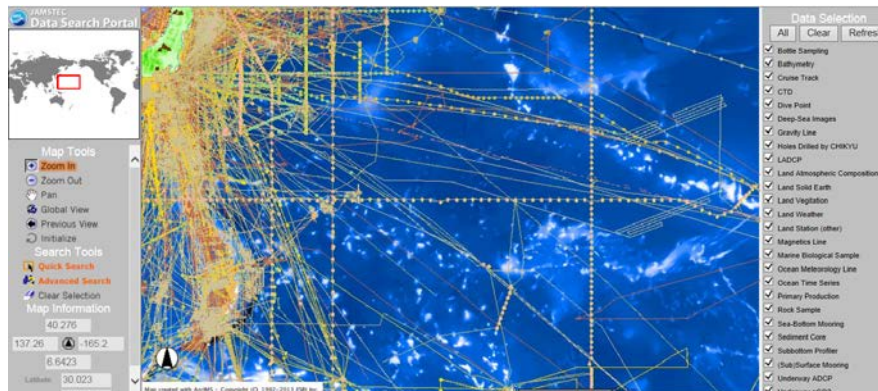
# Good Practice for Cruise Data and Samples

## Metadata may include

- | Location
- | Depth
- | Temperature
- | Salinity
- | pH
- | Oxygen content
- | Seafloor conditions

## Sample storage

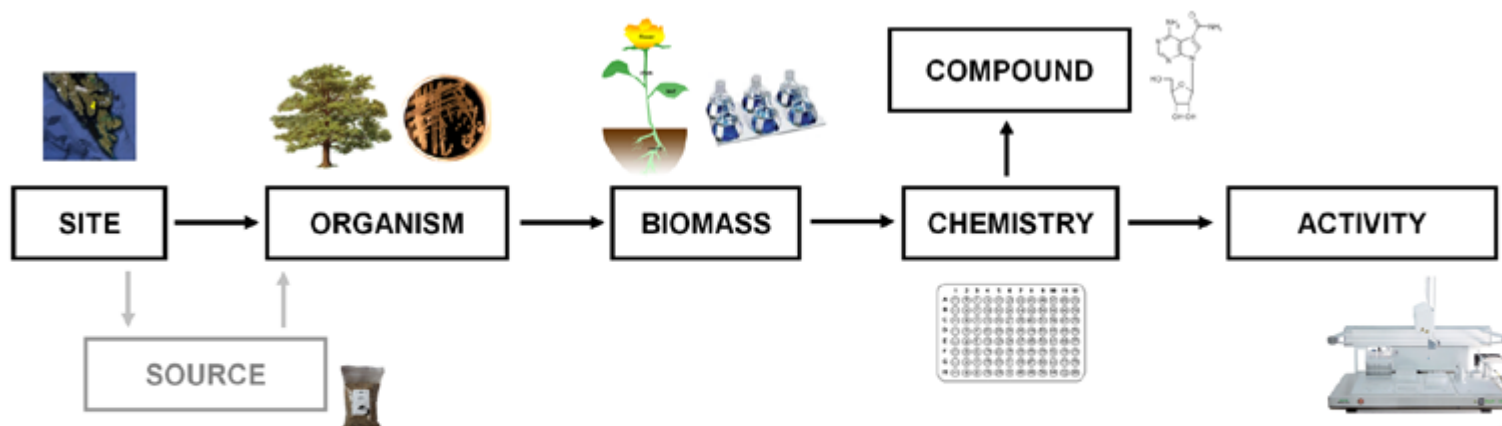
- | Ambient temperature
- | Cooler (4°C)
- | Freezer (-20°C)
- | -80°C Freezer
- | Liquid nitrogen (-196°C)
- | Formaldehyde
- | Ethanol
- | DNA/RNA preservation liquids



**ADD:**  
**Use/Change of use**

# Extract/Pure Compound and Assay Data Management

Possible to track sample from origin to exploitation  
(but better databases are needed)

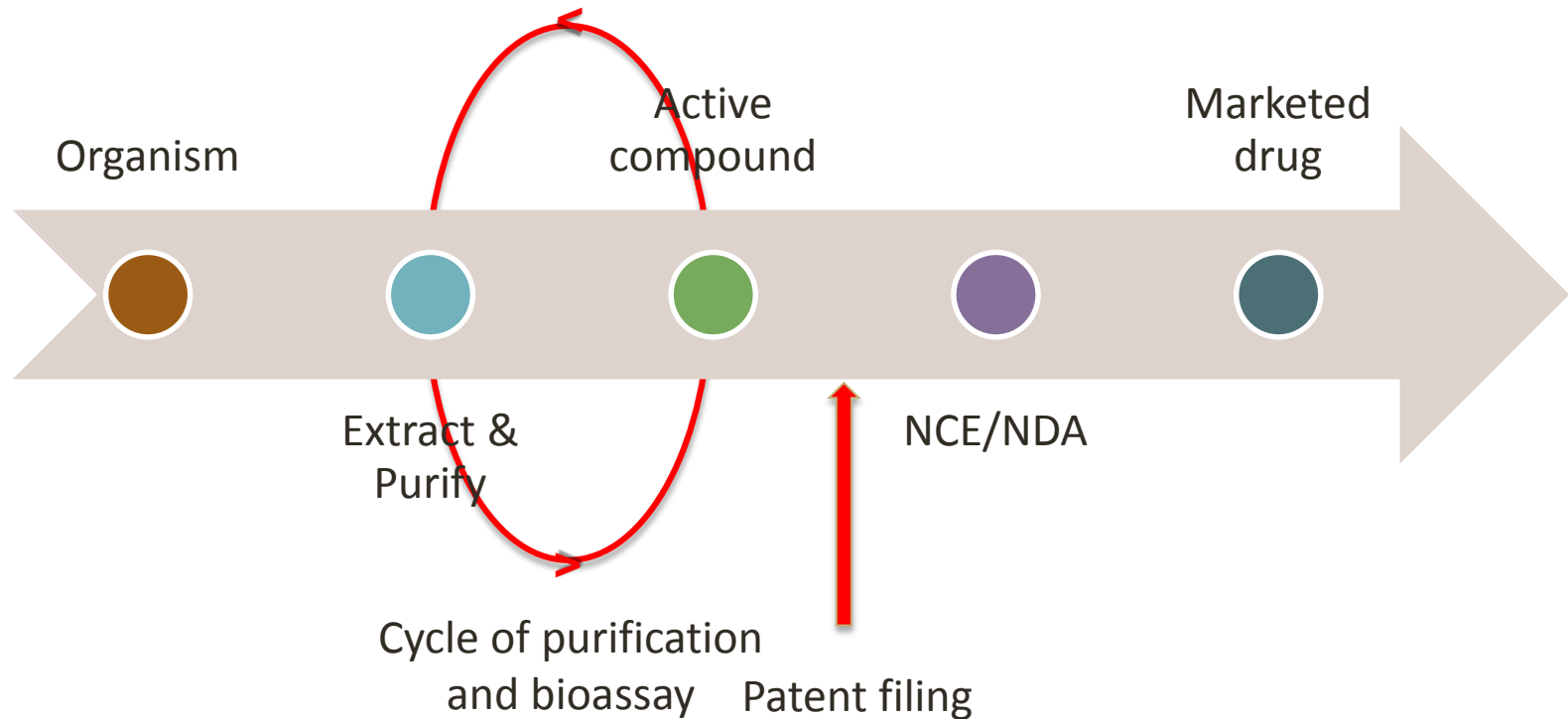


**OpenNAPIS™**  
Functional Design

White Point Systems, Inc.  
20100626

Example of open access system: [www.pangaea.de](http://www.pangaea.de)

# Value Inflection Point Model



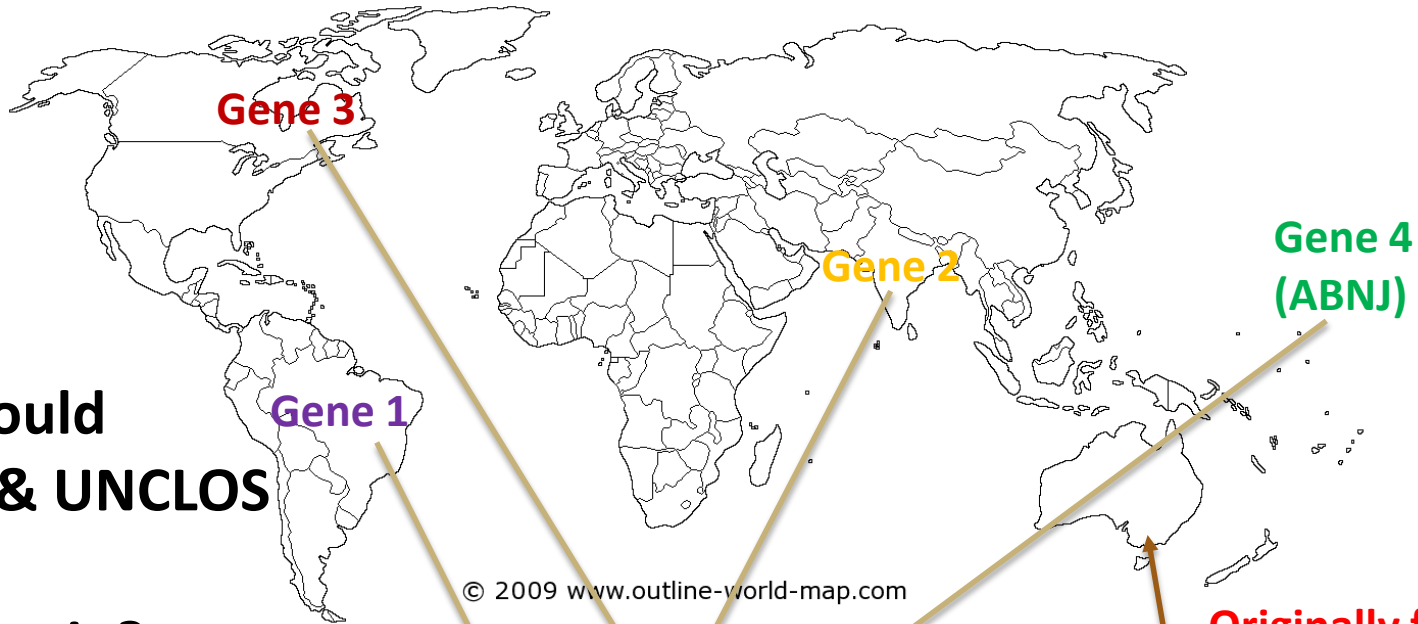
# Value Inflection Point Model

- Where is the transition from basic research to research with commercial intent? (when does actual value become apparent?)
  - When research is initiated (statement in grant application?)
  - When organism is found to be active against disease based screen?
  - When pure compound with activity is identified?
  - When patent is filed?
  - When NCE is found and NDA is filed?
  - When commercial involvement begins? (ie industry funding/licencing of NCE to pharma)
- Not all patents commercialised. Some industries don't patent.
- Collection permits required?
- Who maintains data & how is use monitored?
- How is environmental impact assessed?

# Science is Moving Very Quickly

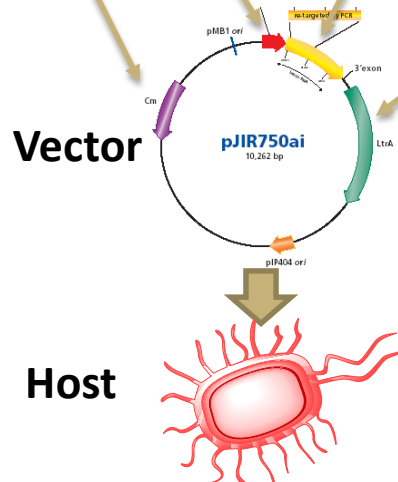
- Gene sequencing is now inexpensive and extremely rapid
  - Sequence whole genomes quickly
  - Sequence metagenomes
- Bioinformatics is making huge advances
  - Genome mining to find enzymes/compounds
  - Understanding of how to express these compounds in alternative hosts
- Low cost gene synthesis is a game-changer
  - No longer need genetic resource, only sequence
  - Sequences can be optimised to express in alternative host
  - Genes can be easily combined
- **But:** open databases don't always indicate origin of materials.
- **Any UNCLOS implementing agreement developed over the next few years may not be flexible enough to deal with rapid scientific progress.**

# Possible Scenario

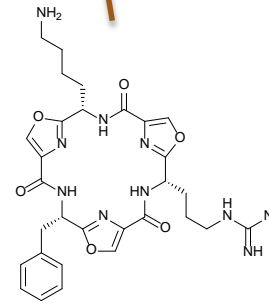


What would Nagoya & UNCLOS do with this scenario?

Vector and host may have associated IP rights



Originally found in Australian EEZ Marine organism



Known bioactive compound

# Possible Scenario

- This is based on the use of synthetic biology in which genes from a number of different organisms are combined to make a product.
- The genes are combined in a ‘vector’ which allows them to work together. Vectors are artificial DNA constructs which are often owned by companies or have associated IP.
- The vector is incorporated into a ‘host’ organism, a modified easy-to-grow microorganism that can express the genes in the vector. The hosts are often owned by companies or have associated IP.
- Whilst in the host, the genes placed within the vector produce the proteins that biosynthesise the small molecules of interest.
- In this scenario the molecule generated was initially isolated from an Australian Great Barrier Reef seasquirt (marine invertebrate).



# Origin of the Genes

- Gene from Brazilian reef organism – cloned without further modifications (actual gene taken from the organism unmodified and inserted into the vector).
- Gene from Indian marine cyanobacterium (blue-green alga). Organism collected by Indian scientist, sequenced, and whole genome deposited online in public database. Gene synthesised without modifications and incorporated in vector.
- Gene from Canadian marine sponge. Genome sequenced and deposited in online public database. Gene synthesised *with major modifications* and incorporated in vector.
- Gene from marine microorganism isolated from sediment obtained from deep sea trench. Trench is in ABNJ. (Gene can be native, synthetic or modified synthetic as in 1-3 above)

# Open Access Model

- Current problems
  - Access is difficult to control and monitor
  - Nagoya requires researcher to prove samples are not from national jurisdiction
  - Synthetic biology will change everything – genes used may be heavily modified and bear no similarity to original genes
- Solution: share samples and data
- Open Access to Biodiversity Beyond National Jurisdiction?
- Open access typically used when:
  - There is no desire/need to control access
  - There is more than enough of a resource (samples/data) for all to utilise

# Open Access Model

- Open Access Precedents
  - Biology - open data resources (also BioBricks)
  - Software – open source software can be used to make profit making products
  - Semiconductor industry – eg all contributors to chip design get right to exploit in their markets.
- Low cost – commensurate with size of problem
- Benefits will accrue locally on exploitation (jobs, knowledge, service industries etc)
- Is current good practice sufficient?
  - Access monitoring
  - Environmental impact
  - Data logging policy (time to deposition?)
  - Need for policing?

# How Can We Ensure All Can Benefit?

- All should be able to benefit from discoveries
- Open access/open source typically leads to greater innovation, transparency and openness
- Requires capacity building to ensure fairness
- Access for landlocked & developing countries
  - Participate in cruises?
  - Access to actual samples (direct sampling strategy)
  - Access to open data - sample metadata, (meta)genomic, proteomic etc (make sure location/organism is clearly stated)
- Make sure all can benefit and can exploit
  - Develop scientific expertise
  - Build scientific infrastructure
  - Help identify/build local markets

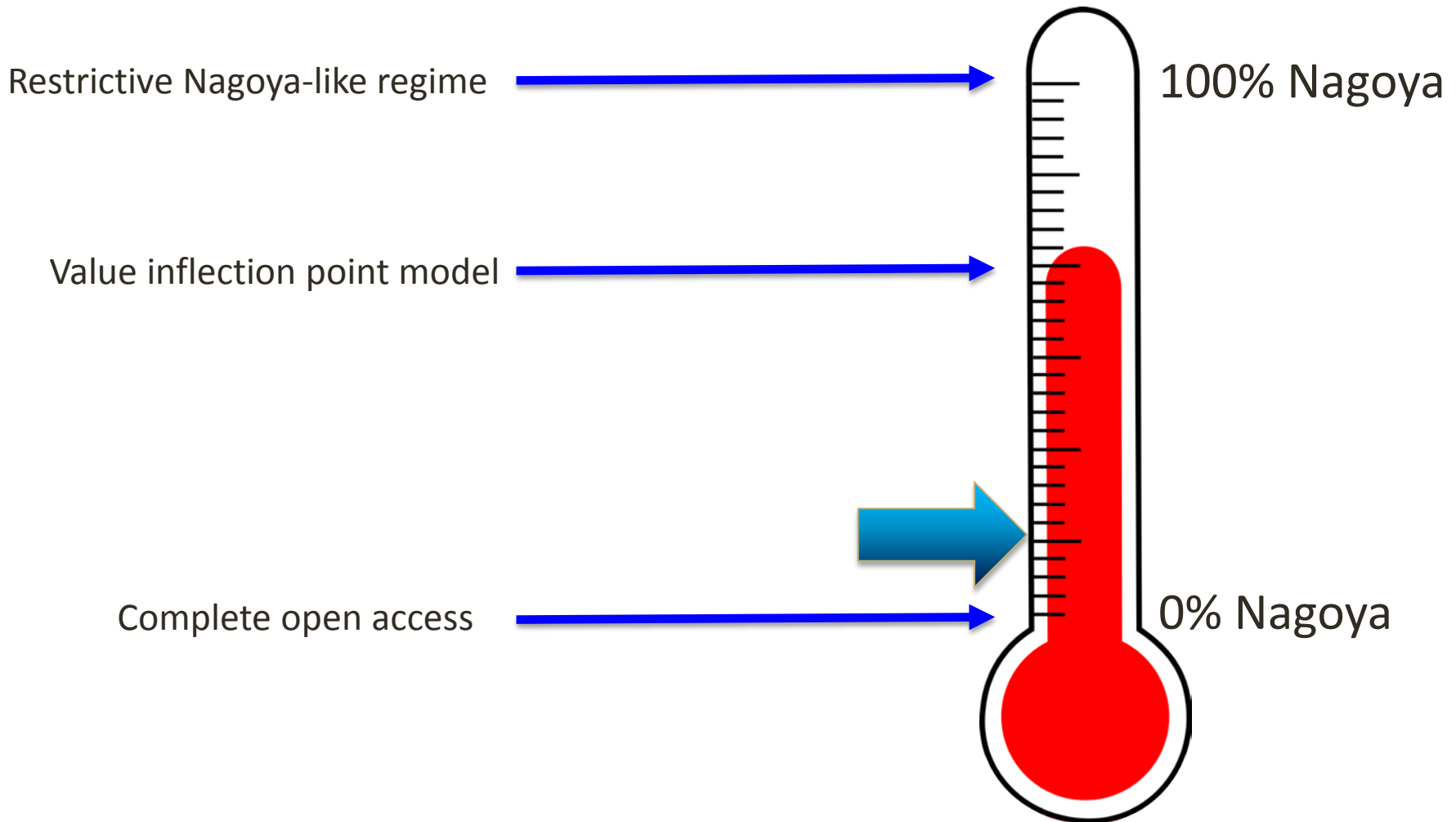
# Aspects of UNCLOS Compatible with Open Access

- Rights of MSR include international cooperation and dissemination/knowledge transfer
- Apply Bermuda principles (rapid deposition of genomic data)
- Main goal of UNCLOS implementing agreement is to protect and conserve biodiversity
- Most MSR is basic research and benefits biodiversity conservation

## **BUT**

- Environmental impact assessment difficult as no baseline data

# Nagoya-O-Meter



# Conclusions

- Nagoya style ABS regime not workable for multiple reasons.
- Value inflection ABS model difficult to implement as there is no defined point where ‘commercial intent’ is declared in all cases.
- Any new UNCLOS implementing agreement will struggle to be flexible enough to accommodate pace of scientific progress
- Pragmatic solution is one that builds on idea of open access:
  - Collection of BBNJ not large scale (gauge importance of BBNJ)
  - Current good practice with minor changes may be sufficient as monitoring tool – much infrastructure already in place
  - Much data already open access and new data likely to be open access too
  - All can utilise/all can benefit
  - Requires capacity building so all can truly benefit
- What kind of legal regime is necessary to make open access workable?

# PHARMASEA



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