



Recent developments in mathematical virology

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Overview

- Virus structure and dynamics
 - Icosahedral symmetry
 - Tiling theory
 - Extended structures
 - Giant viruses
- 2 Virus assembly
 - MS2 and Packaging Signals
 - Virus-like Particles
 - Anti-virals
- 3 Disease dynamics
 - Intracellular Replication
 - Immunological Infection Dynamics
 - Modelling

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What is a Virus?

- Transported piece of genetic information that e.g. can run a programme in a host cell
- Genome: RNA or DNA
- Fragile needs to be protected by a protein shell: capsid
- Gene → mRNA → protein (transcription and translation)
- Each protein = amino acid chain folds into a 3D shape: one geometric building block









Watson and Crick: The Icosahedron







- Crick&Watson: Genetic economy → symmetry → icosahedral is largest
- Rotational icosahedral group is $I = A_5$ of order 60
- Full icosahedral group is the Coxeter group H₃ of order 120 (including reflections/inversion); generated by the root system icosidodecahedron

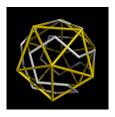
Many viruses are icosahedral



Assembling an Icosahedron







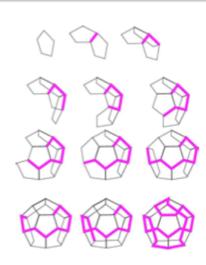
- Assemble from 20 identical triangular building blocks
- The order of addition gives a Hamiltonian path on the dual dodecahedron

Icosahedral symmetry Tiling theory

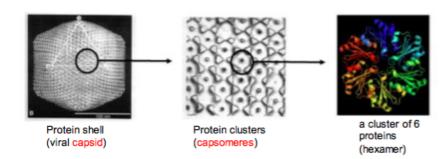
Tiling theory
Extended structures
Giant viruses

Assembling a dodecahedron





More than just icosahedral symmetry?

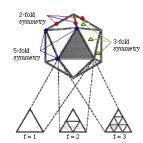


Caspar and Klug: Triangulations

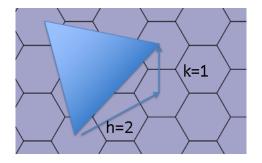
- Mathematical upper limit of 60 for equivalent subunits, but biologically want to do better!
- Gene → can already make a triangle → might as well make many!
- Caspar-Klug ideas of quasi-equivalence and triangulations







Viruses: Caspar-Klug triangulations $T = h^2 + hk + k^2$



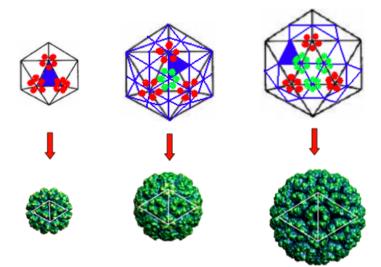
integer steps h and k in hexagonal directions give allowed triangulation numbers $T = h^2 + hk + k^2$

T orbits so $60\,T$ proteins, 60 of which form 12 pentamers, and 60(T-1) form 10(T-1) hexamers



Icosahedral symmetry Tiling theory Extended structures Giant viruses

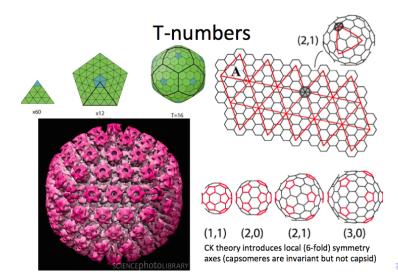
Viruses: Caspar-Klug triangulations $T = h^2 + hk + h^2$





Icosahedral symmetry Tiling theory Extended structures Giant viruses

Viruses: Caspar-Klug triangulations





The status quo for 40 years

 New insights in the 2000s from Reidun Twarock who essentially founded the field and other mathematical physicists (e.g. Anne Taormina etc)



Fullerenes

- other icosahedral objects in nature: football-shaped fullerenes
- Different shells with icosahedral symmetry: e.g. C_{60} , C_{240} , C_{540}
- Follow Caspar-Klug-like layouts (e.g. $T=h^2$ and $T=3h^2$ families)

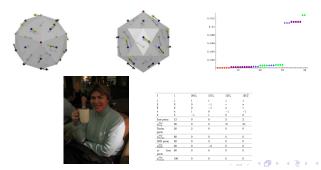






Vibrations of capsids and fullerenes

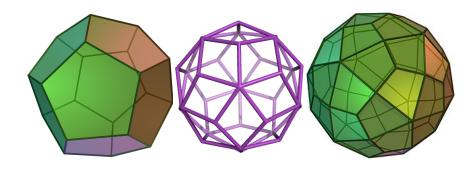
- Normal modes/vibrations of icosahedral capsids given by representation theory of the icosahedral group
- E.g. $\Gamma_{\text{Loos}}^{\text{disp}} = \Gamma^1 + 3\Gamma^3 + \Gamma^{3'} + 2\Gamma^4 + 3\Gamma^5$
- Pioneered by Anne Taormina, Kasper Peeters and Francois Englert



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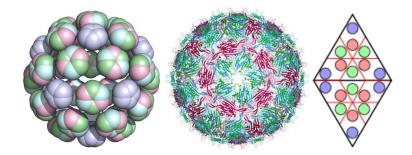
More general icosahedral tilings



Other tile shapes can also give icosahedral tilings: pentagons (dodecahedron), rhombuses (rhombic triacontahedron), kites (deltoidal hexecontahedron)



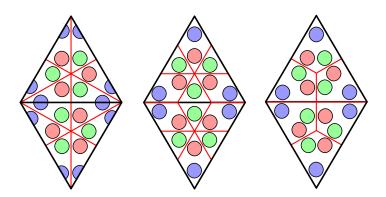
triangulations vs other quasi-equivalent tilings



Two viral surface layouts: a T=4 triangulation (e.g. HBV) and a rhombus tiling (MS2) for a pseudo T=3 triangulation



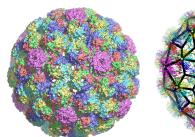
Other quasi-equivalent tilings

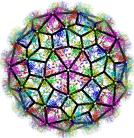


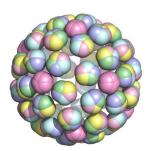
Three T = 3 capsids: Polio, MS2 and Pariacoto



A puzzle: non-quasiequivalent tilings – Penrose







More general icosahedral tilings: Cryo-EM reconstruction of HPV, a kite-rhombus tiling and a pseudo $\mathcal{T}=7$ triangulation.

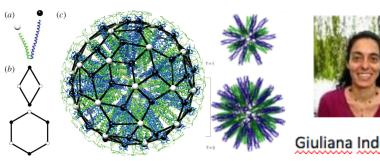


Architecture

- Triangulations: Buckminster Fuller geodesic domes
- kite-rhombus tiling: the new Amazon HQ



Self-assembling protein nanoparticles





Giuliana Indelicato

Quantized e.g. in mass spec - predict units by symmetry. Particles eg. for vaccine design

More general symmetry still?

- Improves the limit to 60 T, but only in terms of surface structures (12 pentagons and rest hexagons).
- Making the symmetry non-compact might allow more general symmetry, simultaneously constraining different 'radial levels'
- Non-compact generator is a translation motivates looking into affine extensions of icosahedral symmetry
- There is an inherent length scale in the problem given by size of nucleic acid/protein molecules

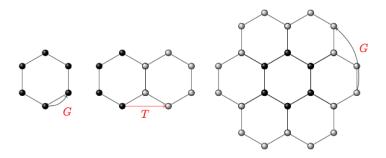


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Affine extensions - A_2

Unit translation along a vertex of a unit hexagon



A random translation would give 6 secondary hexagons, i.e. 36 points. Here we have degeneracies due to 'coinciding points', and building up the hexagonal lattice.

Affine extensions of non-crystallographic groups?

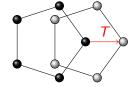
Unit translation along a vertex of a unit pentagon



Affine extensions of non-crystallographic groups?

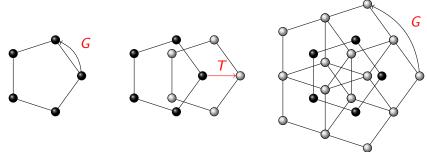
Unit translation along a vertex of a unit pentagon





Affine extensions of non-crystallographic groups?

Unit translation along a vertex of a unit pentagon

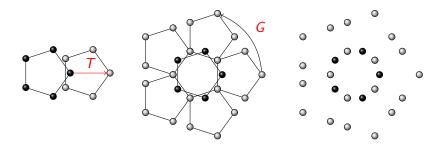


A random translation would give 5 secondary pentagons, i.e. 25 points. Here we have degeneracies due to 'coinciding points'.



Affine extensions of non-crystallographic root systems?

Translation of length $\tau = \frac{1}{2}(1+\sqrt{5}) \approx 1.618$ (golden ratio)

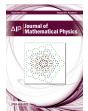


Cartoon version of a virus or carbon onion. Would there be an evolutionary benefit to have more than just compact symmetry?

The problem has an intrinsic length scale.

Affine extensions of non-crystallographic Coxeter groups

- 2D and 3D point arrays for applications to viruses, fullerenes, quasicrystals, proteins etc
- Two complementary ways to construct these



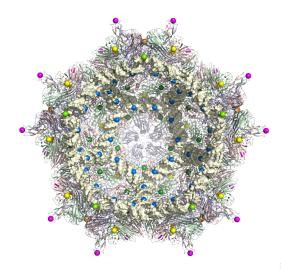




Other ideas

- Project symmetry orbits in 6D to get finite extended icosahedral point arrays (Emilio Zappa)
- Use projection to find 3D tiles to model viruses (David Salthouse)
- Use projection to model transitions between capsids via lattice transitions in 6D (Giuliana Indelicato)

Use in Mathematical Virology



New insight into RNA virus assembly

- There are specific interactions between RNA and coat protein (CP) given by icosahedral symmetry axes
- Essential for assembly, as only this RNA-CP interaction turns
 CP into right geometric shape for capsid formation
- Hamiltonian cycle visiting each RNA-CP contact once dictated by symmetry
- Even the RNA has an icosahedrally ordered component









Extension to fullerenes: carbon onions

- Extend idea of affine symmetry to other icosahedral objects in nature: football-shaped fullerenes (with Jess Wardman)
- Recover different shells with icosahedral symmetry from affine approach: carbon onions $(C_{60}-C_{240}-C_{540})$









Extension to fullerenes: carbon onions

- Extend idea of affine symmetry to other icosahedral objects in nature: football-shaped fullerenes
- Recover different shells with icosahedral symmetry from affine approach: carbon onions $(C_{80} C_{180} C_{320})$

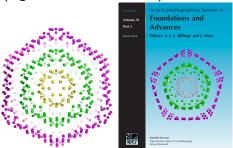


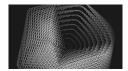




Viruses and fullerenes – symmetry as a common thread?

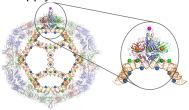
 Get nested arrangements like Russian dolls: carbon onions (e.g. Nature 510, 250253)





Two examples

- Non-compact symmetry that relates different structural features in the same polyhedral object when there is an additional length scale
- Novel symmetry principle in Nature, shown that it seems to apply to at least fullerenes and viruses





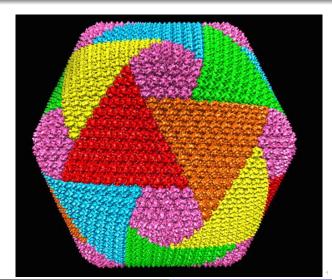




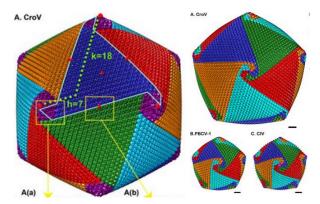
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Giant viruses



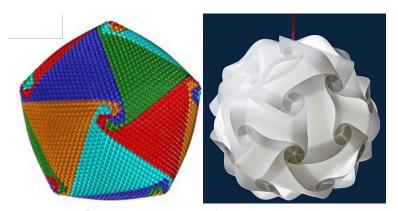
A common approach – little hooks



Pentasymmetrons and trisymmetrons



A common approach – little hooks



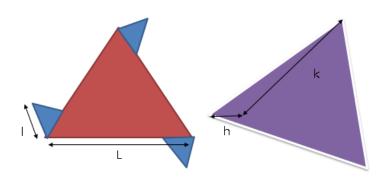
Pentasymmetrons and trisymmetrons



A family of solutions: h = 7 – and some gaps

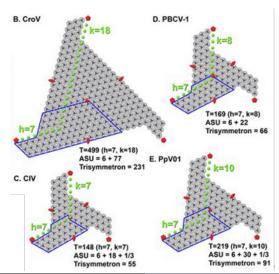
- Chilo iridescent virus: T = 147, h = 7 and k = 7
- Paramecium bursaria Chlorella virus 1: T = 169, h = 7 and k = 8
- Phaeocystis pouchetti virus: T = 219, h = 7 and k = 10
- Faustovirus: T = 277, h = 7 and k = 12
- Pacman virus: T = 309, h = 7 and k = 13
- Cafeteria roenbergensis: T = 499, h = 7 and k = 18

Count areas?

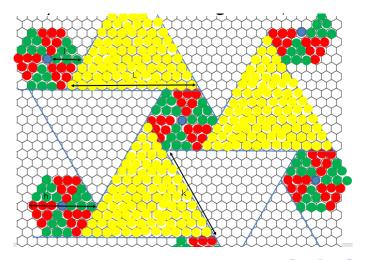


When is there an equivalent description in terms of Caspar-Klug *h* and *k*?

Count hexagons



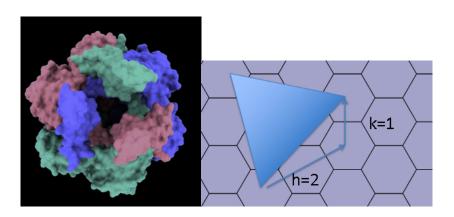
Trisymmetrons and Pentasymmetrons – how to count



Trisymmetrons and Pentasymmetrons – how to count

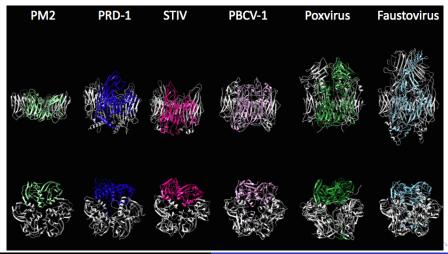
- Each has I(I+1)/2 or L(L+1)/2 hexamers
- So in total 20(3/(l+1)/2 + L(L+1)/2)
- Caspar-Klug theory predicts $10(T-1) = 10(h^2 + hk + k^2 1)$
- So for which h, k, l, L can there be equality?
- Turns out for h = 2l + 1 and k = L l this holds identically
- So can have odd h and any k

Major capsid protein



T is an area, so \sqrt{T} gives size of triangle and thus also particle diameter イロン イ御 とくきと くきと

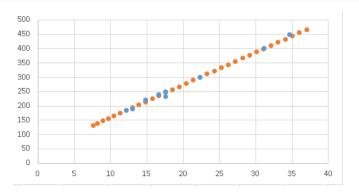
Major capsid protein – evolutionary conservation



Pierre-Philippe Dechant

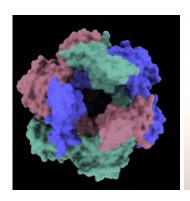
Recent developments in mathematical virology

Scaling



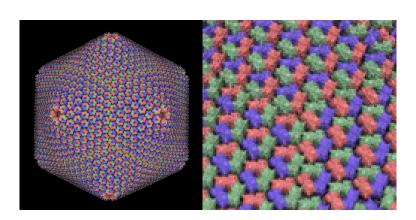
Missing points allowed geometrically but less stable? Or just not yet discovered? Predict Tetraselmis virus 1 TetV-1 of $257nm \pm 9nm$ is exactly T=343. Predict holes in family exist and sizes given by this scaling

Major capsid protein – trimer / pseudohexamer

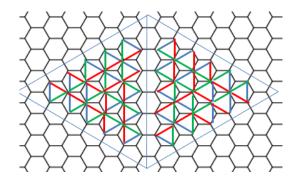




Decorations – threefold and giants



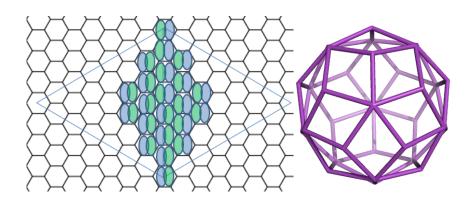
Decorations – threefold and giants



Use hexagonal tiling unit with decorations that partially break the symmetry: 6, 3, 2, 1

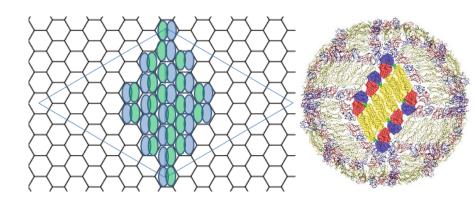
If respect the 3-fold axis, then still have partial lattice symmetry! Defects/domain walls at the other symmetry axes

Decorations – twofold case



Symmetric with respect to 2-fold axis – quasi-lattice symmetry Looks like a rhombic triacontahedron all over again

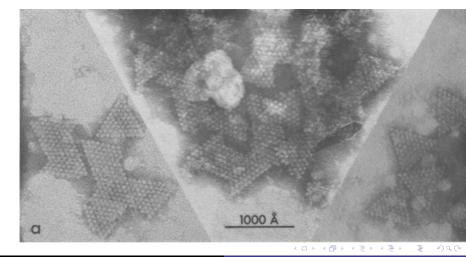
Decorations – twofold and Zika



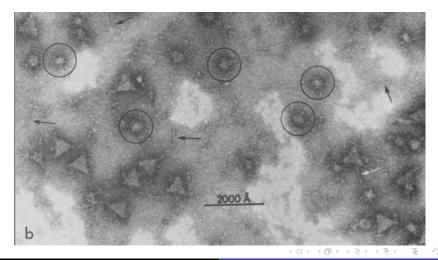
This is exactly what Zika looks like



Trisymmetrons and Pentasymmetrons



Major capsid protein - trimer, pseudohexamer



Build from prearranged blocks? Back to Hamiltonian paths







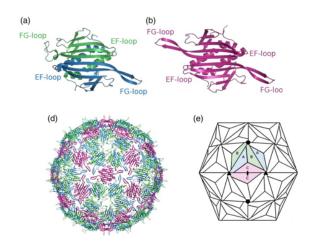
- Are the trisymmetrons and pentasymmetrons preformed? (or is that just what virions fall apart into?)
- If trisymmetrons are assembled then we're back to a Hamiltonian path for the icosahedron
- If pentasymmetrons then get a slightly new polyhedron



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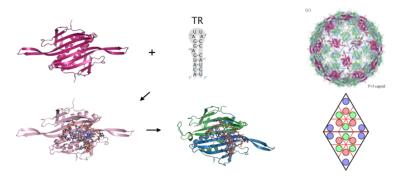
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MS2 tiling and dimeric building blocks: A/B and C/C



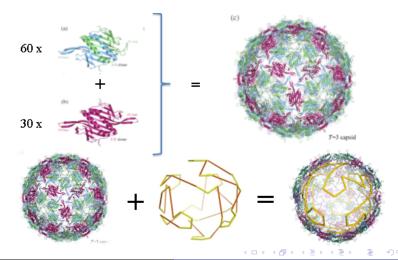
Need to bind RNA in 60 places

The TR sequence is known to initiate assembly by associating with the maturation protein. It forms a <u>stemloop</u> and it has been shown that the <u>stemloop</u> changes the conformation of the <u>symmetric</u> C/C dimer to the <u>asymmetric</u> A/B dimer (allosteric switch).



Peter Stockley (Leeds), Neil Ranson (Leeds), Eric Dykeman (York)

MS2 Hamiltonian path



New insight into RNA virus assembly

- More realistic examples for MS has 60 vertices with 41,000 paths
- The RNA is actually circularised by Maturation Protein: only 66 cycles
- With thermodynamical assembly kinetics and 5-fold averaging experiments uniquely idenfied an evolutionarily conserved cycle
- Patents for new antiviral strategies and virus-like nanoparticles
 e.g. for drug delivery (Twarock group)



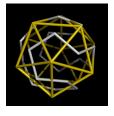






Hamiltonian cycles on icosahedral solids







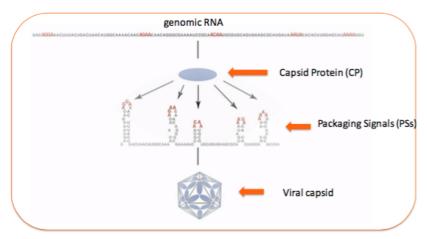
- So interaction contacts are given by the symmetry
- Orbits of the interaction points have to be visited by the RNA exactly once
- Even the RNA has an icosahedrally ordered component
- Hamiltonian cycles for dodecahedron, icosahedron and rhombic triacontahedron



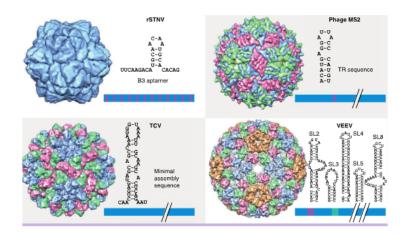
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Multiple dispersed Packaging Signals paradigm



Common Mechanism across groups of viruses



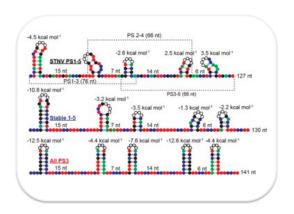
German Leonov, Richard Bingham



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Engineering Packaging Signals to make VLPs

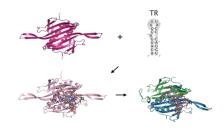


Improved sequences assemble twice as efficiently (Nikesh Patel). Potential applications to vaccines or drug delivery.

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Understanding assembly allows one to interfere



- target RNA
- target CP
- introduces competitors
- this might drive evolution due to exerting selection pressures



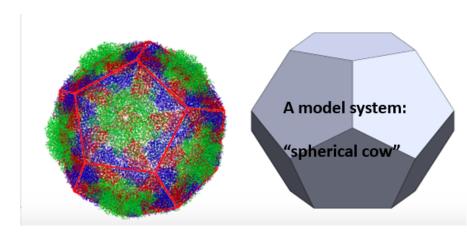
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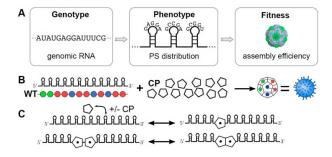
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Dodecahedral cow (Rich Bingham)



Viral evolution and quasispecies

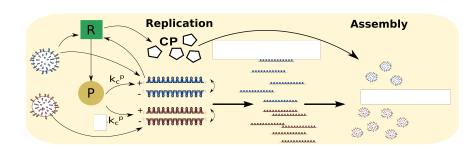


A phenomenological genome space of 12 packaging signals with 3 binding affinity bands (weak, medium, strong). Can compute the whole space explicitly in terms of assembly efficiency.

Simulations

- Stochastic simulations rather than ODE models because of discrete nature and low numbers
- Gillespie algorithm that selects a random reaction to occur (Eric Dykeman)
- Couple an intracellular model (replication) with an infection model (immune system)

Intracellular model: replication



Intracellular reactions

$$p_V^+ + R \stackrel{k_r^{on}}{\longleftrightarrow} (p_V^+, R)$$
 (Ribosome positive strand virus binding/unbinding)

$$(p_V^+,R) \xrightarrow{k_v^c} p_V^+ + R + P$$
 (Genome translation - makes P and abundant CP)

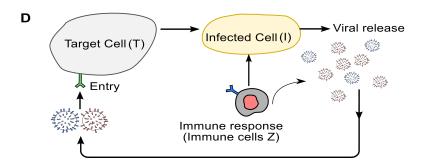
$$p_{V/S}^{\pm} + P \underset{\stackrel{k}{\leftarrow} off}{\overset{k_p^{\pm on}}{\leftarrow}} (p_{V/S}^{\pm}, P)$$
 (Polymerase positive/negative strand virus binding/unbinding)

$$(p_{V/S}^{\pm}, P) \xrightarrow{k_p^c} p_{V/S}^{\pm} + p_{V/S}^{\mp} + P$$
 (complementary strand production)

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 - Modelling

Infection model: immune system



Immune cell reactions

$$T \xrightarrow{\lambda} 2T$$
 (Target cell birth)
$$T \xrightarrow{d_T} 0 \text{ (Target cell death)}$$

$$T + pV + qS \xrightarrow{\beta} I \text{ (Infection of target cell)}$$

$$I \xrightarrow{a} rV + sS \text{ (Infected cell death/lysis, if } p > 0)$$

$$I + Z \xrightarrow{\pi} Z \text{ (Infected cell removal by immune system)}$$

$$V + Z \xrightarrow{u} Z \text{ (Virion removal by immune system)}$$

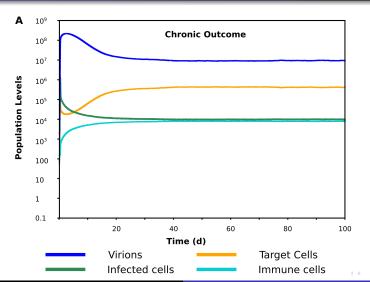
$$I + Z \xrightarrow{c} I + 2Z \text{ (Immune cell birth)}$$

$$Z \xrightarrow{b} 0 \text{ (Immune cell death)}$$

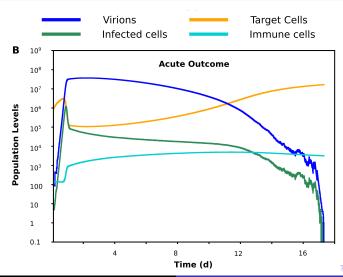
Overview

- Virus structure and dynamics
 - Icosahedral symmetry
 - Tiling theory
 - Extended structures
 - Giant viruses
- 2 Virus assembly
 - MS2 and Packaging Signals
 - Virus-like Particles
 - Anti-virals
- 3 Disease dynamics
 - Intracellular Replication
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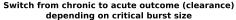
Chronic infections

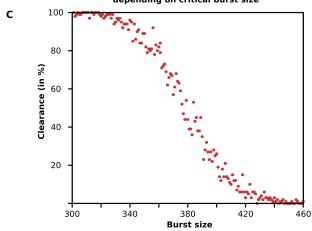


Acute infections

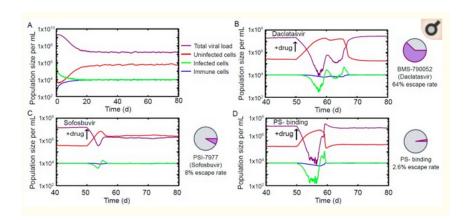


Transition between chronic and acute





Evolutionarily stable drugs



Summary of mathematical virology

- a truly diverse interdisciplinary endeavour: group theory, tiling theory, dynamical systems, graph theory, computational modelling, biophysics, bioinformatics, biochemistry, cell biology, structural biology, immunology
- curiosity-driven and impactful: from understanding a structural puzzle to a new generation of anti-virals – by accident

Other algebraic interests

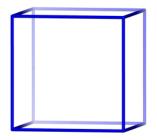
- exceptional root systems/geometries
- (reflexive) polytopes
- Clifford algebras
- ADE correspondences

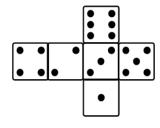
Intracellular – Replication Immunological – Infection Dynamics Modelling

Thank you! LGBTQ+ lunch at 1pm.

Symmetry averaging

- As an example use the symmetry of a cube
- (axes of 4-fold, 3-fold and 2-fold symmetry)
- Dice have approximately cubic symmetry, but have slightly asymmetric features (the faces)



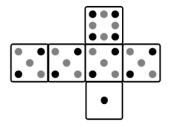


Symmetry averaging

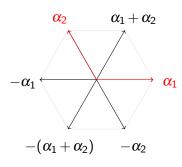
- As an example use the symmetry of a cube
- Approximately cubic symmetry, but have slightly asymmetric features (the faces)
- Experiment averages over all orientations
- Washes out all asymmetric features

Symmetry averaging

- Assume we have a distinguished face, e.g. 1 is always at the bottom
- The 6 can be in two configurations
- Average over the other 4 faces



Root systems



Root system Φ : set of vectors α in a vector space with an inner product such that

1.
$$\Phi \cap \mathbb{R}\alpha = \{-\alpha, \alpha\} \ \forall \ \alpha \in \Phi$$

$$2. s_{\alpha} \Phi = \Phi \ \forall \ \alpha \in \Phi$$

Simple roots: express every element of Φ via a \mathbb{Z} -linear combination.

reflection/Coxeter groups
$$s_{\alpha}: v \to s_{\alpha}(v) = v - 2 \frac{(v|\alpha)}{(\alpha|\alpha)} \alpha$$

Affine extensions

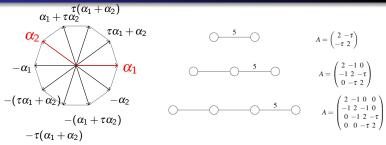
An affine Coxeter group is the extension of a Coxeter group by an affine reflection in a hyperplane not containing the origin $s_{\alpha_0}^{aff}$ whose geometric action is given by

$$s_{\alpha_0}^{aff} v = \alpha_0 + v - \frac{2(\alpha_0|v)}{(\alpha_0|\alpha_0)} \alpha_0$$

Non-distance preserving: includes the translation generator

$$Tv = v + \alpha_0 = s_{\alpha_0}^{aff} s_{\alpha_0} v$$

Non-crystallographic Coxeter groups $H_2 \subset H_3 \subset H_4$



 $H_2 \subset H_3 \subset H_4$: 10, 120, 14,400 elements, the only Coxeter groups that generate rotational symmetries of order 5 linear combinations now in the extended integer ring

$$\boxed{\mathbb{Z}[\tau] = \{a + \tau b | a, b \in \mathbb{Z}\}} \text{ golden ratio} \boxed{\tau = \frac{1}{2}(1 + \sqrt{5}) = 2\cos\frac{\pi}{5}}$$

$$\boxed{x^2 = x + 1} \boxed{\tau' = \sigma = \frac{1}{2}(1 - \sqrt{5}) = 2\cos\frac{2\pi}{5}} \boxed{\tau + \sigma = 1, \tau\sigma = -1}$$