Group Actions Applied to Virus Architecture

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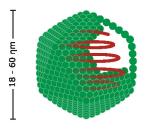
joint work with A. Devillers (UWA) and R. Twarock (Univ. of York, UK)

Workshop on Symmetry in Graphs, Maps and Polytopes

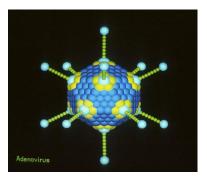
October 25th, 2011 = 200th anniversary of Galois!!!

Fields Institute, Toronto, 24–27 October 2011

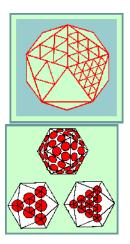
Viruses



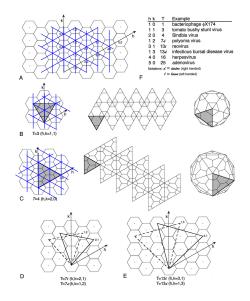
- 1956 Crick and Watson: "The only reasonable way to build a protein shell is to use the same type of molecule over and over again." "subunits would be packed so as to provide each with an identical environment. This is possible only if they are packed symmetrically." They predicted *cubic* symmetry.
- Later: icosahedral symmetry is preferred in virus structure.



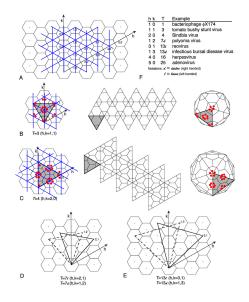
- subunits of few different types in *quasi-equivalent* environments.
- triangulations of icosahedron faces with capsomers in corners of triangles.



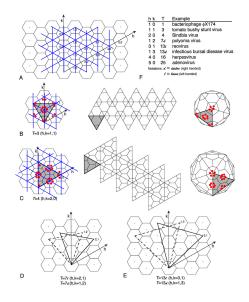
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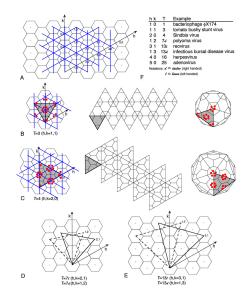
- (*h*, *k*) are co-ordinates of vertex of icosahedron.
- embedding of icsahedron in triagular lattice.
- A capsomer in corner of each triangle.
- T number is square of distance between two icosahedron vertices. This is exactly h² + hk + k².
- There are 60T capsomers, clustered as 12 pentamers and 10(T - 1) hexamers.



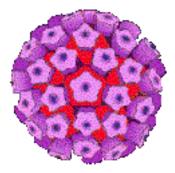
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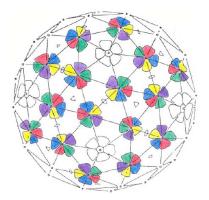
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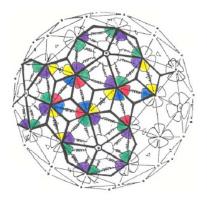
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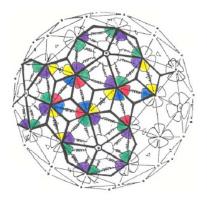
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- T = 7
- Twarock 2004: Use other tiles!
- Capsomers only at corners which have same angle
- Rhombs and kites tiling also explains dimer and trimer interactions.



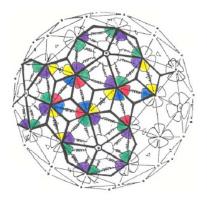
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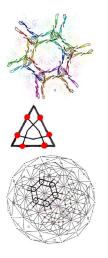


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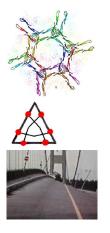
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Viral Tiling Theory



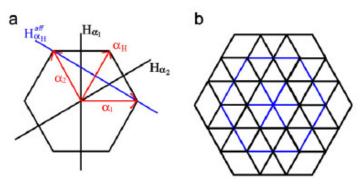
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- Also dynamical properties of the capsid. How genetic material is released; resonant frequencies of the capsid

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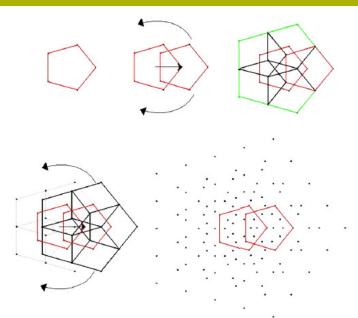
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- *crystallographic groups* stabilize a lattice (i.e. a Z-span of a basis).
- Coxeter groups of type H₃ and H₄ are not crystallographic.
- *H*₃ is the icosahedral group.
- There are "lattice-like" structures...



We extend the group $sl(3,\mathbb{C})$ with a translation. This yields a triangular lattice.

Affine extension of H_2

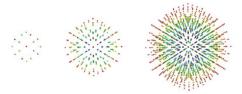


Philippe Cara (VUB)

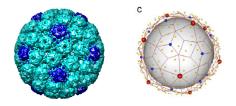
- We extend the group C₅ with a translation T. This yields a dense orbit.
- If we restrict the number of times we are allowed to use the translation, we get part of quasi-lattice.
- For $s \in \mathbb{N}$ we define $M^s = \{$ words made from $C_5 \cup \{T\}$ where T appears at most s times $\}$.

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Layers in capsid structure



(Keef, Twarock, Elsawy 2008)

• Consider a finite group $G \leq GL(\mathbb{R}^k)$. Example: \mathcal{I} .

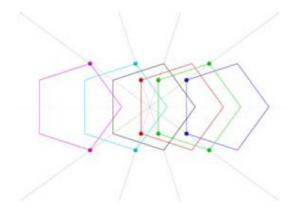
- For v ∈ ℝ^k we can look at the orbit G(v), a starting configuration.
 Example: the 12 vertices of an icosahedron.
- Locate the axes of (rotational) symmetry of G(v).
- A translation T along an axis of symmetry is called admissible if there are at least two elements $u_1, u_2 \in G(v)$ for which Tu_1 and Tu_2 both on (different) axes of symmetry.

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Admissible translations for C_5



	five fold	three fold	two fold
v vertexv center of facev center of edge	3	4	6
	4	3	5
	6	5	5

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v center of face	4	3	5
v center of edge	6	5	5

 $M^1 = \mathcal{I} \cup \{g_1 T g_2 \mid g_1, g_2 \in \mathcal{I}\}$

• VIPER database contains data about capsids of many viruses.

- Choose a virus.
- Compare outer layer to 41 outer layers we have.
- Scale the whole model.
- For Paricoto virus we get a good prediction of the fine structure of the capsid.
- You can combine several clouds.
- Group theoretical X-ray analysis of viruses!

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