# Algebraic Geometry in Applications

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Lahore, August 2018

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# SINGULAR



### A Computer Algebra System for Polynomial Computations

with special emphasize on the needs of algebraic geometry, commutative algebra, and singularity theory

W. Decker, G.-M. Greuel, G. Pfister, H. Schönemann Technische Universität Kaiserslautern Fachbereich Mathematik; Zentrum für Computer Algebra D-67663 Kaiserslautern

http://www.mathematik.uni-kl.de/ pfister/vortragLahore.pdf

### Elimination

#### lexikographical ordering

 $x_1^{\alpha_1} \cdot \ldots \cdot x_n^{\alpha_n} > x_1^{\beta_1} \cdot \ldots \cdot x_n^{\beta_n}$  if  $\alpha_j = \beta_j$  for  $j \le k-1$  and  $\alpha_k > \beta_k$ 

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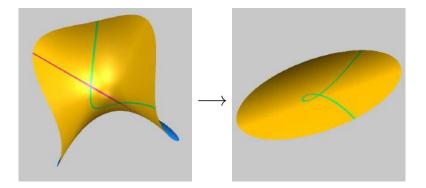
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 I ⊂ K[x<sub>1</sub>,...,x] ideal, G Gröbner basis, then G ∩ K[x<sub>k</sub>,...,x<sub>n</sub>] is a Gröbner basis of I ∩ K[x<sub>k</sub>,...,x<sub>n</sub>].
 this means geometrically to compute the projection π : V(I) ⊂ K<sup>n</sup> → K<sup>n-k+1</sup>.

### Projection



# $\pi: V(z^2-x+1,y-xz) \subset \mathbb{C}^3 \longrightarrow V(x^3-x^2-y^2) \subset \mathbb{C}^2.$

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# Infineon Tricore Project



- Aim: prove that the processor (32 Bit) works correctly
- every instruction of the processor will be verified specifying special properties and proving them
- it is difficult to check the arithmetic properties

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Algebraic Geometry in Applications

### Robotics

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## Roboti<u>cs</u>

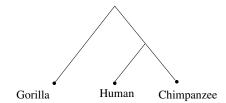


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Phylogenetics is the study of the evolution of a set of species from a common ancestor. The evolution will be described using a **phylogenetic tree**.



To reconstruct such a tree pieces of DNA sequences are used.

	ヨン・ハロン・ハロン
Human	AAGCTTCACCGGCGCAGTCATTCTCATAATCGCCCACGGGCTTACATCCT
Cimpanzee	AAGCTTCACCGGCGCAATTATCCTCATAATCGCCCACGGACTTACATCCT
Gorilla	AAGCTTCACCGGCGCAGTTGTTCTTATAATTGCCCACGGACTTACATCAT

## **Computer Vision**



The Perspective-n-Point problem, i.e. the problem of determining the absolute position and orientation of a camera given its intrinsic parameters and a set of n 2D-to-3D point correspondences, is one of the most important problems in computer vision with a broad range of applications in structure from motion or recognition. Felix Kubler and Karl Schmedders (University of Zürich)

General problem:

Study a computer model of a national economy,

a standard exchange economy with finitely many agents and goods

especially study equilibria

Walrasian equilibrium consists of prices and choices, such that household maximize utilities, firms maximize

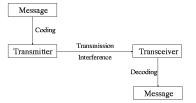
profits and markets clear

Mathematical problem:

Find the positive real roots of a given system of polynomial equations

# Coding theorie





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Sudoku

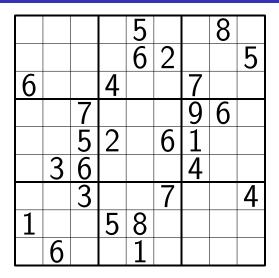


Abbildung: Sudoku

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# A Problem of Group Theory Solved Using Algebraic Gometry and Computer Algebra

Let G be a finite group, define

$$G^{(1)} := [G,G] = \langle aba^{-1}b^{-1} \mid a,b \in G \rangle.$$

and  $G^{(i)} := [G^{(i-1)}, G]$ . G is called nilpotent, if  $G^{(m)} = \{e\}$  for some m.

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- Abelian groups are nilpotent.
- If the order of G is a power of a prime, G is nilpotent.
- G is nilpotent  $\Leftrightarrow$  it is a direct product of its Sylow groups.
- S<sub>3</sub> is not nilpotent.

## Nilpotent Groups

Magma:

- > G:=Sym(3);
- > H:=CommutatorSubgroup(G,G);

Η;

Permutation group acting on a set of cardinality 3

Order = 3

(1, 2, 3)

```
> CommutatorSubgroup(H,G);
```

Permutation group acting on a set of cardinality 3

Order = 3

(1, 2, 3)

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### Dihedral Group

$$D_4 = < r, s | r^4 = s^2 = e, srs = r^{-1} >$$

> #DihedralGroup(4);

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```
> G:=CommutatorSubgroup(DihedralGroup(4),DihedralGroup(4));
```

Permutation group acting on a set of cardinality 4

Order = 2

(1, 3)(2, 4)

> CommutatorSubgroup(G,DihedralGroup(4));

Permutation group acting on a set of cardinality 4

Order = 1

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Now define

$$G^{(i)} := [G^{(i-1)}, G^{(i-1)}],$$

then G is called solvable, if  $G^{(m)} = \{e\}$  for a suitable m.

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- nilpotente groups are solvable.
- $S_3, S_4$  are solvable.
- groups of odd order are solvable.
- $S_5, A_5$  are not solvable.



$$\mathsf{PSL}(2, K) = \left. \mathsf{SL}(2, K) / \left\{ \left( \begin{smallmatrix} a & 0 \\ 0 & a \end{smallmatrix} \right) \; \middle| \; a^2 = 1 \right\} \right.$$

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especially

$$\begin{aligned} \mathsf{PSL}(2,\mathbb{F}_5) &= \{ \left[ \begin{pmatrix} a_{11} & a_{12} \\ a_{21} & a_{22} \end{pmatrix} \right], \ a_{11}a_{22} - a_{21}a_{12} = 1 \} \\ & \left[ \begin{pmatrix} a_{11} & a_{12} \\ a_{21} & a_{22} \end{pmatrix} \right] &= \left\{ \begin{pmatrix} a_{11} & a_{12} \\ a_{21} & a_{22} \end{pmatrix}, \ \begin{pmatrix} 4a_{11} & 4a_{12} \\ 4a_{21} & 4a_{22} \end{pmatrix} \right\}. \end{aligned}$$

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$$\mathsf{PSL}(2, K) = \left. \mathsf{SL}(2, K) / \left\{ \left(\begin{smallmatrix} a & 0 \\ 0 & a \end{smallmatrix}\right) \right| a^2 = 1 \right\}$$

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$$\mathsf{PSL}(2, \mathbb{F}_5) = \{ \left[ \begin{pmatrix} a_{11} & a_{12} \\ a_{21} & a_{22} \end{pmatrix} \right], \ a_{11}a_{22} - a_{21}a_{12} = 1 \} \\ \left[ \begin{pmatrix} a_{11} & a_{12} \\ a_{21} & a_{22} \end{pmatrix} \right] = \left\{ \begin{pmatrix} a_{11} & a_{12} \\ a_{21} & a_{22} \end{pmatrix}, \ \begin{pmatrix} 4a_{11} & 4a_{12} \\ 4a_{21} & 4a_{22} \end{pmatrix} \right\}.$$

It holds:

$$\mathsf{PSL}(2,\mathbb{F}_5)\cong \mathsf{PSL}(2,\mathbb{F}_4)\cong A_5$$

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## Solvable groups

```
> G:=PSL(2,5);
> G;
Permutation group G acting on a set of cardinality 6
Order = 60 = 2^2 * 3 * 5
   (3, 4)(5, 6)
   (1, 6, 2)(3, 4, 5)
> IsIsomorphic(G,Alt(5));
true Homomorphism of GrpPerm: G, Degree 6, Order 2^2 * 3 * 5 into
GrpPerm: $, Degree 5, Order 2^2 * 3 * 5 induced by
   (3, 4)(5, 6) \mid --> (1, 3)(2, 5)
   (1, 6, 2)(3, 4, 5) \mid --> (1, 4, 2)
```

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**Problem**: Characterize the class of finite solvable groups *G* by 2-variable identities.

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Example:

• *G* is abelian  $\Leftrightarrow xy = yx \ \forall x, y \in G$ 

• (Zorn, 1930) A finite group G is **nilpotent**  $\Leftrightarrow \exists n \ge 1$ , such that  $v_n(x, y) = 1 \forall x, y \in G$ (Engel Identity)

**Theorem** (T. Bandman, G.-M. Greuel, F. Grunewald, B. Kunyavsky, G. Pfister, E. Plotkin)

 $U_1 = U_1(x, y) := x^{-2}y^{-1}x,$  $U_{n+1} = U_{n+1}(x, y) = [xU_nx^{-1}, yU_ny^{-1}].$ 

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A finite group G is **solvable**  $\Leftrightarrow \exists n$ , such that  $U_n(x, y) = 1 \forall x, y \in G$ .

• Let  $x, y \in G$  such that  $y \neq x^{-1}$  and  $U_1(x, y) = U_2(x, y) \Rightarrow U_n(x, y) \neq 1 \forall n \in \mathbb{N}.$ 

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G solvable  $\Rightarrow$  Identity is true (by definition).

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Idea of  $\Leftarrow$ 

**Theorem** (Thompson, 1968) Let G be simple and minimally not solvable (i.e. G is not solvable but every proper subgroup is solvable). Then G is one of the following groups:

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• **PSL** $(2, \mathbb{F}_p)$ , *p* a prime number  $\geq 5$ 

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- **PSL** $(2, \mathbb{F}_{2^p})$ , *p* a prime number
- **PSL** $(2, \mathbb{F}_{3^p})$ , *p* a prime number

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■ **PSL**(3, **F**<sub>3</sub>)

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- **Sz** $(2^p)$  *p* a prime number.

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- **PSL**(3, **F**<sub>3</sub>)
- **Sz** $(2^p)$  *p* a prime number.

If is enough to prove (for *G* in Thompson's list):

 $\exists x, y \in G$ , such that  $y \neq x^{-1}$  and  $U_1(x, y) = U_2(x, y)$ .

# Translation to algebraic Geometry

Let us consider 
$$G = \mathsf{PSL}(2, \mathbb{F}_p), p \ge 5$$

Image: Image:

Let us consider 
$$G = \mathsf{PSL}(2, \mathbb{F}_p), p \ge 5$$

Consider the matrices

$$x = \left( egin{array}{c} t & 1 \ -1 & 0 \end{array} 
ight) \qquad y = \left( egin{array}{c} 1 & b \ c & 1+bc \end{array} 
ight)$$

 $x^{-1} = \left( egin{smallmatrix} 0 & -1 \ 1 & t \end{smallmatrix} 
ight)$  implies  $y 
eq x^{-1}$  for all  $(b,c,t) \in \mathbb{F}_p^3$ .

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$$U_1(x, y) = U_2(x, y)$$
, i.e.  
 $x^{-1}yx^{-1}y^{-1}x^2 = yx^{-2}y^{-1}xy^{-1}$ 

has a solution  $(b, c, t) \in \mathbb{F}_p^3$ .

### The equations

The entries of  $U_1(x, y) - U_2(x, y)$  are the following polynomials  $p_1, \ldots, p_4$  in  $\mathbb{Z}[b, c, t]$ . Let  $I = \langle p_1, \ldots, p_4 \rangle$ .

$$\begin{split} p_1 &= b^3 c^2 t^2 + b^2 c^2 t^3 - b^2 c^2 t^2 - bc^2 t^3 - b^3 ct + b^2 c^2 t + b^2 ct^2 + 2bc^2 t^2 \\ &+ bct^3 + b^2 c^2 + b^2 ct + bc^2 t - bct^2 - c^2 t^2 - ct^3 - b^2 t + bct + c^2 t \\ &+ ct^2 + 2bc + c^2 + bt + ^2 ct + c + 1 \end{split} \\ p_2 &= -b^3 ct^2 - b^2 ct^3 + b^2 c^2 t + bc^2 t^2 + b^3 t - b^2 ct - 2bct^2 - b^2 c + bct \\ &+ c^2 t + ct^2 - bt - ct - b - c - 1 \cr p_3 &= b^3 c^3 t^2 + b^2 c^3 t + b^2 c^2 t + b^2 ct^2 + b^3 c^2 t + b^2 c^3 t + b^2 c^2 t^2 \\ &+ bc^2 t + ct^2 + bct^2 + b^2 c^2 t + b^2 ct^2 + bc^2 t^2 - c^2 t^3 - ct^4 - 2b^2 ct \\ &+ bc^2 t + c^3 t + bct^2 + 2c^2 t^2 + ct^3 - b^2 c - b^2 t + bct + c^2 t + bt^2 \\ &+ 3ct^2 + bc - bt - b - c + 1 \cr p_4 &= -b^3 c^2 t^2 - b^2 ct^2 + ct^3 + b^2 t - bct - c^2 t - ct^2 + b^2 - bt \\ &- 2ct - b - t + 1 \end{split}$$

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#### The zero set of I is a curve.

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The entries of  $U_1(x, y) - U_2(x, y)$  are the following polynomials  $p_1, \ldots, p_4$  in  $\mathbb{Z}[b, c, t]$ . Let  $I = \langle p_1, \ldots, p_4 \rangle$ .

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#### The zero set of I is a curve.

We have to prove that for every prime p there are  $\mathbb{F}_p$ -rational points on the curve.

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# **Theorem von Hasse–Weil** (generalized by Aubry and Perret for singulare curves):

**Theorem von Hasse–Weil** (generalized by Aubry and Perret for singulare curves): Let  $C \subseteq \mathbb{A}^n$  be an absolutely irreducible affine curve defined over

the finite field  $\mathbb{F}_q$  and  $\overline{C} \subset \mathbb{P}^n$  its projective closure  $\Rightarrow$ 

 $\#C(\mathbb{F}_q) \ge q + 1 - 2p_a\sqrt{q} - d$ 

 $(d = \text{degree}, p_a = \text{arithmetic genus of } \overline{C}).$ 

**Theorem von Hasse–Weil** (generalized by Aubry and Perret for singulare curves):

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The Hilbert-polynomial of  $\overline{C}$ ,  $H(t) = d \cdot t - p_a + 1$ , can be computed using the ideal  $I_h$  of  $\overline{C}$ : We obtain  $H(t) = 10t - 11 \Rightarrow d = 10$ ,  $p_a = 12$ . **Theorem von Hasse–Weil** (generalized by Aubry and Perret for singulare curves):

Let  $C \subseteq \mathbb{A}^n$  be an absolutely irreducible affine curve defined over the finite field  $\mathbb{F}_q$  and  $\overline{C} \subset \mathbb{P}^n$  its projective closure  $\Rightarrow$ 

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The Hilbert-polynomial of  $\overline{C}$ ,  $H(t) = d \cdot t - p_a + 1$ , can be computed using the ideal  $I_h$  of  $\overline{C}$ : We obtain  $H(t) = 10t - 11 \Rightarrow d = 10$ ,  $p_a = 12$ . Since  $p + 1 - 24\sqrt{p} - 10 > 0$  if p > 593, we obtain the result.

### **Proposition:** $V(I^{(p)})$ is absolutely irreducibel for all primes $p \ge 5$ .

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Using **SINGULAR** we show:

 $\langle f_1, f_2 \rangle : h^2 = I.$ 

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Using **SINGULAR** we show:

$$\langle f_1, f_2 \rangle : h^2 = I.$$

$$\begin{split} f_1 &= t^2 b^4 + (t^4 - 2t^3 - 2t^2) b^3 - (t^5 - 2t^4 - t^2 - 2t - 1) b^2 \\ &- (t^5 - 4t^4 + t^3 + 6t^2 + 2t) b + (t^4 - 4t^3 + 2t^2 + 4t + 1) \\ f_2 &= (t^3 - 2t^2 - t) c + t^2 b^3 + (t^4 - 2t^3 - 2t^2) b^2 \\ &- (t^5 - 2t^4 - t^2 - 2t - 1) b - (t^5 - 4t^4 + t^3 + 6t^2 + 2t) \\ h &= t^3 - 2t^2 - t \end{split}$$

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### Let $P(x) := t^2 J[1]|_{b=x/t}$ then P is monic of degree 4.

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Let  $P(x) := t^2 J[1]|_{b=x/t}$  then P is monic of degree 4.

$$\begin{aligned} & x^4 + (t^3 - 2t^2 - 2t)x^3 - (t^5 - 2t^4 - t^2 - 2t - 1)x^2 - \\ & (t^6 - 4t^5 + t^4 + 6t^3 + 2t^2)x + (t^6 - 4t^5 + 2t^4 + 4t^3 + t^2). \end{aligned}$$

We prove, that the induced polynomial  $P \in \mathbb{F}_p[t, x]$  is absolutely irreducibel for all primes  $p \ge 2$ . (Using the lemma of Gauß this is equivalent to P being irreducibel in  $\overline{\mathbb{F}}_p(t)[x]$ .) Ansatz

(\*) 
$$P = (x^2 + ax + b)(x^2 + gx + d)$$

a, b, g, d polynomials in t with variable coefficients

a(i), b(i), g(i), d(i).

Ansatz

(\*) 
$$P = (x^2 + ax + b)(x^2 + gx + d)$$

a, b, g, d polynomials in t with variable coefficients

a(i), b(i), g(i), d(i).

The decomposition (\*) with a(i), b(i), g(i), d(i)  $\in \overline{\mathbb{F}}_p$  does not exist iff the ideal C generated by the coefficients with respect to x, t of  $P - (x^2 + ax + b)(x^2 + gx + d)$  has no solution in  $\overline{\mathbb{F}}_p$ . This is equivalent to the fact that  $1 \in \mathbb{C}$ .

## The ideal of the coefficients of C:

```
C[1] = -b(5) * d(3)
C[2] = -b(5) * g(2)
C[3] = -b(4)*d(3)-b(5)*d(2)
C[4] = -b(4)*g(2)-b(5)*g(1)-d(3)-1
C[5] = -b(3)*d(3)-b(4)*d(2)-b(5)*d(1)+1
C[6] = -b(5) - g(2) - 1
C[7] = a(0)*b(5)-a(2)*d(3)-b(3)*g(2)-b(4)*g(1)-d(2)+4
C[8] = -a(0)^{2}+b(5)+b(0)+b(5)-b(2)+d(3)-b(3)+d(2)-b(4)+d(1)-b(5)-4
C[9] = -a(2) * g(2) - b(4) - g(1) + 2
C[10] = a(0)*b(4)-a(1)*d(3)-a(2)*d(2)-b(2)*g(2)-b(3)*g(1)-d(1)-1
C[11] = -a(0)^{2} + b(4) + b(0) + b(4) - b(1) + d(3) - b(2) + d(2) - b(3) + d(1) - b(4) + 2
C[12]=a(0)-a(1)*g(2)-a(2)*g(1)-b(3)-d(3)
C[13] = -a(0)^{2}+a(0)*b(3)-a(0)*d(3)-a(1)*d(2)-a(2)*d(1)+b(0)-b(1)*g(2)-b(2)*g(1)-7
C[14] = -a(0)^{2}b(3)+b(0)*b(3)-b(0)*d(3)-b(1)*d(2)-b(2)*d(1)-b(3)+4
C[15] = -a(2) - g(2) - 2
C[16]=a(0)*a(2)-a(0)*g(2)-a(1)*g(1)-b(2)-d(2)+1
C[18] = -a(0)^{2*b(2)+b(0)*b(2)-b(0)*d(2)-b(1)*d(1)-b(2)+1}
C[19] = -a(1) - g(1) - 2
C[20] = a(0)*a(1)-a(0)*g(1)-b(1)-d(1)+2
C[21] = -a(0)^{2}a(1)+a(0)*b(1)-a(0)*d(1)+a(1)*b(0)-a(1)-b(0)*g(1)
C[22] = -a(0)^{2}*b(1)+b(0)*b(1)-b(0)*d(1)-b(1)
C[23] = -a(0)^{3}+2*a(0)*b(0)-a(0)
C[24] = -a(0)^{2*b}(0) + b(0)^{2-b}(0)
                                                             ▲口 ▶ ▲冊 ▶ ▲目 ▶ ▲目 ▶ ● ● ● ●
```

#### Using $\operatorname{SINGULAR}$ , one shows that over

# $\mathbb{Z}[\{a(i)\}, \{b(i)\}, \{g(i)\}, \{d(i)\}]$

$$4=\sum_{i=1}^{24}M_i\,\,\mathrm{C}[i]\,.$$

Gerhard Pfister

### Algebraic Statistics: Point of View of Algebraic Geometry

#### A statistical model in algebraic statistics is a polynomial map

$$\varphi: \mathbb{R}^d \to \mathbb{R}^m$$

#### Example

If X is the random variable <sup>1</sup> describing the number of heads in m flips of a coin, and  $t \in [0, 1]$  is the probability that we obtain head in one flip, then we can use the **binomial distribution** to model this situation:

$$\operatorname{Prob}(X=j) = \binom{m}{j} t^j (1-t)^{m-j} .$$

 $^1A$  random variable is defined as a function that maps the outcomes of unpredictable processes to numerical quantities, typically real numbers.  $\equiv$ 

**Gerhard Pfister** 

Algebraic Geometry in Applications

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These polynomials describe a map

$$\varphi: \mathbb{R} \to \mathbb{R}^{m+1}, t \mapsto (\dots, {m \choose i} t^j (1-t)^{m-j}, \dots).$$

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Algebraic Geometry in Applications

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#### This map is our statistical model.

 $^{1}$ A random variable is defined as a function that maps the outcomes of unpredictable processes to numerical quantities, typically real numbers.  $\geq$ 

Gerhard Pfister

#### Example

If we regard  $\varphi$  as a map over the complex numbers

$$\varphi: \mathbb{C} \to \mathbb{C}^{m+1}, t \mapsto (\dots, {m \choose j} t^j (1-t)^{m-j}, \dots),$$

write  $p_0, \ldots, p_m$  for the coordinate functions on  $\mathbb{C}^{m+1}$ , and consider the ideal

$$J := \langle \{p_j - \binom{m}{j} t^j (1-t)^{m-j} \}_{j=0,\dots,m} \rangle \subseteq \mathbb{C}[p_0,\dots,p_m,t],$$

then the elimination ideal  $I = J \cap \mathbb{C}[p_0, \dots, p_m]$  describes the Zariski closure of the image of  $\varphi$ .

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Gerhard Pfister

Based on observations in an experiment, we can use the model invariants to value *t*. We consider the case m = 6:

#### Example

#### Example

```
I[1] = p(0)+p(1)+p(2)+p(3)+p(4)+p(5)+p(6)-1
```

```
I[2] =5*p(5)^2-12*p(4)*p(6)
```

```
...
I[16]=5*p(1)^2+7560*p(1)*p(6)+12600*p(2)*p(6)
+16200*p(3)*p(6)+18900*p(4)*p(6)+21000*p(5)*p(6)
+22680*p(6)^2-12*p(2)+54*p(3)-252*p(4)+1680*p(5)
-22680*p(6)
```

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#### Example

Now suppose that we observed in an experiment that  $p_3 = \frac{1}{4}$ . Then this determines the other  $p_i$  in the model.

```
LIB "solve.lib";
I = I, p(3)-1/4;
```

solve(I);

We obtain 6 solutions, 2 of which are real:

Example		
[1]:	[2]:	
[1]:	[1]:	
0.064862202	0.0024089531	
[2]:	[2]:	
0.22479279	0.025023044	
[3]:	[3]:	
0.32460995	0.10830306	
[4]:	[4]:	
0.25	0.25	
[5]:	[5]:	
0.10830306	0.32460995	
[6]:	[6]:	
0.025023044	0.22479279	
[7]:	[7]:	
0.0024089531	0.064862202	¢,

#### Example

From this we deduce that t is either 0.36613231 or 0.63386769. This shows that the coin is not fair (that is, the probability for head is different from  $\frac{1}{2}$ ).

For the general situation let X be a discrete random <sup>2</sup> variable taking values in  $\{1, \ldots, n\}$ . Let the probabilities P(X = i) be given parametrically by polynomials  $p_i(t_1, \ldots, t_d)$ . The statistical model in algebraic statistics is the polynomial map

$$\varphi: \mathbb{R}^d \to \mathbb{R}^n , \varphi(t) = (p_1(t), \ldots, p_n(t)).$$

Consider in  $\mathbb{C}[p_1, \ldots, p_n, t_1, \ldots, t_d]$  the ideal J generated by  $\{p_i - p_i(t)\}_{i=1,\ldots,n}$ . Over the complex numbers the elimination ideal  $I = J \cap \mathbb{C}[p_1, \ldots, p_n]$  describes the Zariski closure of the image of  $\varphi$ , the model variety. Every polynomial in I is called a **model invariant**.

<sup>&</sup>lt;sup>2</sup>A random variable is defined as a function that maps the outcomes of unpredictable processes to numerical quantities, typically real numbers. = +

# Computational Biology

#### What is DNA = Deoxyribo Nucleic Acid?

 DNA molecules contain the biological instructions that make each species unique.

# Computational Biology

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- DNA is made of chemical building blocks called nucleotides. These building blocks are made of three parts: a phosphate group, a sugar group and one of four types of nitrogen bases. To form a strand of DNA, nucleotides are linked into chains.

#### What is DNA = Deoxyribo Nucleic Acid?

- DNA molecules contain the biological instructions that make each species unique.
- DNA is made of chemical building blocks called nucleotides. These building blocks are made of three parts: a phosphate group, a sugar group and one of four types of nitrogen bases. To form a strand of DNA, nucleotides are linked into chains.
- The four types of nitrogen bases found in nucleotides are: adenine (A), thymine (T), guanine (G) and cytosine (C). The order, or sequence, of these bases determines what biological instructions are contained in a strand of DNA.

#### What is DNA = Deoxyribo Nucleic Acid?

- DNA molecules contain the biological instructions that make each species unique.
- DNA is made of chemical building blocks called nucleotides. These building blocks are made of three parts: a phosphate group, a sugar group and one of four types of nitrogen bases. To form a strand of DNA, nucleotides are linked into chains.
- The four types of nitrogen bases found in nucleotides are: adenine (A), thymine (T), guanine (G) and cytosine (C). The order, or sequence, of these bases determines what biological instructions are contained in a strand of DNA.
- DNA contains the instructions needed for an organism to develop, survive and reproduce.

### **Evolution and Mutations**

Evolution depends on **mutations**, that is, changes in the nucleotide sequence of an organisms genetic material.

Given (parts of) the DNA of a number of living species, the goal in using phylogenetic trees is to obtain information on the least common ancestor. The living species are represented by the leaves of the tree, while the root will represent the least common ancestor of all considered species. We make the assumption, that the living species are represented by parts of DNA of equal lenghts:

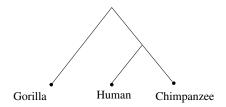
 Gorilla
 AAGCTTCACCGGCGCAGTTGTTCTTATAATTGCCCACGGACTTACATCAT

 Cimpanzee
 AAGCTTCACCGGCGCAATTATCCTCATAATCGCCCACGGACTTACATCCT

 Human
 AAGCTTCACCGGCGCAGTCATTCTCATAATCGCCCACGGGCTTACATCCT

These are strings in the letters A, C, G, T representing the nucleotides.

### Phylogenetic Tree



This tree has 5 nodes including the three leaves corresponding to Gorilla, Human, Chimpanzee.

The node on top of the tree is called root (common ancestor).

We assume that only substitutions occur during the evolutionary process and that this satisfies the following conditions:

- (1) Each nucleotide of the sequence evolves independently of the other nucleotides and in the same way (identically distributed).
- (2) The state at a node only depends on the previous state.<sup>3</sup>
- (3) At bifurcating branches the process is independent on the common node.

<sup>3</sup>A process with this property is called a Markov Process

We consider a so-called phylogenetic tree  $\mathcal{T}$  to model the situation: We think of the edges of the tree as evolutionary steps. We consider a so-called phylogenetic tree  $\mathcal{T}$  to model the situation: We think of the edges of the tree as evolutionary steps. For a node v, we denote by  $P_X^v$  the probability having  $X \in \{A, C, G, T\}$  at a certain position of the DNA string at this node, and write  $P^v = (P_A^v, P_C^v, P_G^v, P_T^v)$ . We consider a so-called phylogenetic tree  $\mathcal{T}$  to model the situation: We think of the edges of the tree as evolutionary steps. For a node v, we denote by  $P_X^v$  the probability having  $X \in \{A, C, G, T\}$  at a certain position of the DNA string at this node, and write  $P^v = (P_A^v, P_C^v, P_G^v, P_T^v)$ . To each edge  $e = (v_1, v_2)$  we associate a matrix of probabilities

$$M_e = \begin{pmatrix} P_{A|A} & \cdots & P_{T|A} \\ P_{A|C} & \cdots & P_{T|C} \\ P_{A|G} & \cdots & P_{T|G} \\ P_{A|T} & \cdots & P_{T|T} \end{pmatrix} = (M(X, Y)),$$

 $M(X, Y) = P_{X|Y}$  is the probability that  $X \in \{A, C, G, T\}$  at the node  $v_1$  changes to  $Y \in \{A, C, G, T\}$  at the node  $v_2$  during the evolutionionary step represented by *e*.  $M_e$  is a **stochastical matrix**.<sup>4</sup>

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 $M(X, Y) = P_{X|Y}$  is the probability that  $X \in \{A, C, G, T\}$  at the node  $v_1$  changes to  $Y \in \{A, C, G, T\}$  at the node  $v_2$  during the evolutionionary step represented by *e*.  $M_e$  is a **stochastical matrix**.<sup>4</sup>

We have  $P^{\nu_1}M_e = P^{\nu_2}$ .

We write  $v_1 = pa(v_2)$  and call  $v_1$  the **parent** of  $v_2$ .

<sup>4</sup>The sum of the entries in a row of the matrix is 1, the sum of the entries of a column in the matrix is 1.

We write for the nodes  $\mathcal{N}(\mathcal{T}) = \{1, \ldots, n, n+1, \ldots, N\}$ such that the leaves  $\mathcal{L}(\mathcal{T}) = \{1, \ldots, n\}$  and N being the root. We assume that we have random variables  $X_1, \ldots, X_N$  at the nodes taking values  $x_1, \ldots, x_N \in \{A, C, G, T\}$  and write

$$P_{x_1,\ldots,x_n} = \operatorname{Prob}(X_1 = x_1,\ldots,X_n = x_n).$$

 Gorilla
 AAGCTTCACCGGCGCAGTTGTTCTTATAATTGCCCACGGACTTACATCAT

 Cimpanzee
 AAGCTTCACCGGCGCAATTATCCTCATAATCGCCCACGGACTTACATCCT

 Human
 AAGCTTCACCGGCGCAGTCATTCTCATAATCGCCCACGGGCTTACATCCT

$$P_{A,A,A} = \frac{\text{number of observations of AAA}}{\text{sequence length}} = \frac{10}{50} = \frac{1}{5}.$$

According to the Markov property of our process we obtain

$$P_{x_1,\ldots,x_n} = \sum_{\substack{(x_{n+1},\ldots,x_N)\\x_s \in \{A,C,G,T\}}} P_{x_N}^N \prod_{v \in \mathcal{N}(\mathcal{T}) \smallsetminus \{N\}} M_{(pa(v),v)}(x_{pa(v)},x_v) \ .$$

According to the Markov property of our process we obtain

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We obtain a map

$$\begin{split} \varphi_{\mathcal{T}} &: \mathbb{R}^{4} \times \prod_{e \in \mathcal{E}(\mathcal{T})} \mathbb{R}^{16} \longrightarrow \mathbb{R}^{4^{n}} \\ \varphi_{\mathcal{T}}(P^{N}, \text{ (entries of } M_{e})_{e \in \mathcal{E}(\mathcal{T})}) = (\dots, P_{x_{1}, \dots, x_{n}}, \dots) \end{split}$$

which we consider as before as a map over the complex numbers:

$$\varphi_{\mathcal{T}}: \mathbb{C}^4 \times \prod_{e \in \mathcal{E}(\mathcal{T})} \mathbb{C}^{16} \longrightarrow \mathbb{C}^{4^n}$$
.

The choice of a special type of the matrices  $M_e$  and a distribution  $P^N$  for the root defines the model  $\mathcal{M}$  choosen for the tree. If these matrices depend on d parameters and  $\pi : \mathbb{C}^d \longrightarrow \mathbb{C}^4 \times \prod_{e \in \mathcal{E}(\mathcal{T})} \mathbb{C}^{16}$ 

defines this specification, we obtain the model map  $\varphi_{\mathcal{T}}^{\mathcal{M}} = \varphi_{\mathcal{T}} \circ \pi$ :

$$\varphi_{\mathcal{T}}^{\mathcal{M}}:\mathbb{C}^{d}\longrightarrow\mathbb{C}^{4^{n}}$$
.

The **phylogenetic variety** according to the tree  $\mathcal{T}$  and the model  $\mathcal{M}$ ,  $V_{\mathcal{M}}(\mathcal{T})$  is the Zariski closure of the image of  $\varphi_{\mathcal{T}}^{\mathcal{M}}$  in  $\mathbb{C}^{4^n}$ .

There are many special models in evolutionary biology. We will give one example. The distribution at the root is usually choosen as  $P^N = \{\frac{1}{4}, \frac{1}{4}, \frac{1}{4}, \frac{1}{4}, \frac{1}{4}\}$ .

There are many special models in evolutionary biology. We will give one example. The distribution at the root is usually choosen as  $P^N = \{\frac{1}{4}, \frac{1}{4}, \frac{1}{4}, \frac{1}{4}, \frac{1}{4}\}$ .

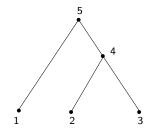
## The Jukes–Cantor model

considers at the edges matrices of type

$$\left( \begin{array}{cccccccc} 1-3a & a & a & a \\ a & 1-3a & a & a \\ a & a & 1-3a & a \\ a & a & a & 1-3a \end{array} 
ight)$$



Let us consider the phylogenetic tree



such that the leaves 1, 2, 3 correspond to Gorilla, Human, Chimpanzee. The the Jukes-Cantor model is given by the map

$$\varphi_T^{\mathcal{M}}: \mathbb{C}^4 \longrightarrow \mathbb{C}^{64}$$

We want to compute now the ideal J of model invariants,  $V(J) = \overline{\Im(\varphi_T^M)}.$ 

We simplify the notations assuming that  $\{A, C, G, T\}$  is identified with  $\{1, 2, 3, 4\}$ . Then we have (before specializing to the Jukes-Cantor model)

$$P_{ijk} = \sum_{l,m=1}^{4} P_m^5 M_{(5,1)}(i,m) M_{(5,4)}(l,m) M_{(4,2)}(j,l) M_{(4,3)}(k,l) .$$

## Example

> ring JC=0,(p(1..4)(1..4)(1..4),a(1..4)),lp;

We create the ideal I associated to the map  $\varphi_T^{\mathcal{M}}$  and eliminate the variables a(1), a(2), a(3), a(4) occuring in the 4 stochastic matrices  $M_{(5,1)}, M_{(5,4)}, M_{(4,2)}, M_{(4,3)}$  to obtian the ideal J of the 61 model invariants.

```
> ideal J=eliminate(I,a(1)*a(2)*a(3)*a(4));
> J;
J[1]=p(4)(4)(2)-p(4)(4)(3)
J[2]=p(4)(4)(1)-p(4)(4)(3)
[...]
```

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3

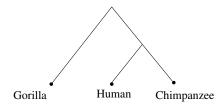
If we compare the parts of DNA for Gorilla, Human and Chimpanzee (from a part of length 1000), we observe  $p_{1,1,1} = \frac{9}{50}$ ,  $p_{4,3,3} = \frac{9}{500}$  and  $p_{1,1,3} = \frac{3}{1000}$ .

If we compare the parts of DNA for Gorilla, Human and Chimpanzee (from a part of length 1000), we observe  $p_{1,1,1} = \frac{9}{50}$ ,  $p_{4,3,3} = \frac{9}{500}$  and  $p_{1,1,3} = \frac{3}{1000}$ . Using this observation we can compute the stochastic matrices. There is one degree of freedom with respect to the 4 parameters of the matrices.

If we put  $a_1 = 0.03$  we obtain  $a_2 = 0.006$ ,  $a_3 = 0.02$  and  $a_4 = 0.05$ .

## Example

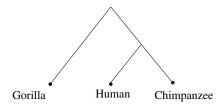
We can use the model invariants to decide about the topology of the tree. In our special situation we have 6 possibilities for Gorilla , Human and Chimpanzee.



We know than this tree is correct.

## Example

We can use the model invariants to decide about the topology of the tree. In our special situation we have 6 possibilities for Gorilla , Human and Chimpanzee.



We know than this tree is correct.

If we exchange the Chimpanzee and the Gorilla in our model then we obtain a value for  $p_{4,1,1} = \frac{3}{1000}$  which was not observed. Observed was  $p_{4,1,1} = \frac{9}{500}$ .